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(54) Title: NOVEL NUCLEIC ACIDS AND POLYPEPTIDES

(57) Abstract:

# NOVEL NUCLEIC ACIDS AND POLYPEPTIDES

## 1. TECHNICAL FIELD

The present invention provides novel polynucleotides and proteins encoded by such polynucleotides, along with uses for these polynucleotides and proteins, for example in therapeutic, diagnostic and research methods.

## 2. BACKGROUND

Technology aimed at the discovery of protein factors (including *e.g.*, cytokines, such as lymphokines, interferons, CSFs, chemokines, and interleukins) has matured rapidly over the past decade. The now routine hybridization cloning and expression cloning techniques clone novel polynucleotides "directly" in the sense that they rely on information directly related to the discovered protein (*i.e.*, partial DNA/amino acid sequence of the protein in the case of hybridization cloning; activity of the protein in the case of expression cloning). More recent "indirect" cloning techniques such as signal sequence cloning, which isolates DNA sequences based on the presence of a now well-recognized secretory leader sequence motif, as well as various PCR-based or low stringency hybridization-based cloning techniques, have advanced the state of the art by making available large numbers of DNA/amino acid sequences for proteins that are known to have biological activity, for example, by virtue of their secreted nature in the case of leader sequence cloning, by virtue of their cell or tissue source in the case of PCR-based techniques, or by virtue of structural similarity to other genes of known biological activity.

Identified polynucleotide and polypeptide sequences have numerous applications in, for example, diagnostics, forensics, gene mapping; identification of mutations responsible for genetic disorders or other traits, to assess biodiversity, and to produce many other types of data and products dependent on DNA and amino acid sequences.

## 3. SUMMARY OF THE INVENTION

The compositions of the present invention include novel isolated polypeptides, novel isolated polynucleotides encoding such polypeptides, including recombinant DNA molecules, cloned genes or degenerate variants thereof, especially naturally occurring variants such as allelic variants, antisense polynucleotide molecules, and antibodies that specifically recognize one or more epitopes present on such polypeptides, as well as hybridomas producing such antibodies.

The compositions of the present invention additionally include vectors, including expression vectors, containing the polynucleotides of the invention, cells genetically engineered to contain such polynucleotides and cells genetically engineered to express such polynucleotides.

The present invention relates to a collection or library of at least one novel nucleic acid sequence assembled from expressed sequence tags (ESTs) isolated mainly by sequencing by hybridization (SBH), and in some cases, sequences obtained from one or more public databases. The invention relates also to the proteins encoded by such polynucleotides, along with therapeutic, diagnostic and research utilities for these polynucleotides and proteins. These nucleic acid sequences are designated as SEQ ID NO: 1-1009. The polypeptides sequences are designated SEQ ID NO: 1010-2018. The nucleic acids and polypeptides are provided in the Sequence Listing. In the nucleic acids provided in the Sequence Listing, A is adenosine; C is cytosine; G is guanine; T is thymine; and N is any of the four bases. In the amino acids provided in the Sequence Listing, * corresponds to the stop codon.

The nucleic acid sequences of the present invention also include, nucleic acid sequences that hybridize to the complement of SEQ ID NO:1-1009 under stringent hybridization conditions; nucleic acid sequences which are allelic variants or species homologues of any of the nucleic acid sequences recited above, or nucleic acid sequences that encode a peptide comprising a specific domain or truncation of the peptides encoded by SEQ ID NO:1-1009. A polynucleotide comprising a nucleotide sequence having at least 90% identity to an identifying sequence of SEQ ID NO:1-1009 or a degenerate variant or fragment thereof. The identifying sequence can be 100 base pairs in length.

The nucleic acid sequences of the present invention also include the sequence information from the nucleic acid sequences of SEQ ID NO:1-1009. The sequence information can be a segment of any one of SEQ ID NO:1-1009 that uniquely identifies or represents the sequence information of SEQ ID NO:1-1009.

A collection as used in this application can be a collection of only one polynucleotide. The collection of sequence information or identifying information of each sequence can be provided on a nucleic acid array. In one embodiment, segments of sequence information is provided on a nucleic acid array to detect the polynucleotide that contains the segment. The array can be designed to detect full-match or mismatch to the polynucleotide that contains the segment. The collection can also be provided in a computer-readable format.

This invention also includes the reverse or direct complement of any of the nucleic acid sequences recited above; cloning or expression vectors containing the nucleic acid sequences; and host cells or organisms transformed with these expression vectors. Nucleic acid sequences (or their reverse or direct complements) according to the invention have numerous applications in a variety of techniques known to those skilled in the art of molecular biology, such as use as hybridization probes, use as primers for PCR, use in an array, use in computer-readable media, use in sequencing

full-length genes, use for chromosome and gene mapping, use in the recombinant production of protein, and use in the generation of anti-sense DNA or RNA, their chemical analogs and the like.

In a preferred embodiment, the nucleic acid sequences of SEQ ID NO:1-1009 or novel segments or parts of the nucleic acids of the invention are used as primers in expression assays that are well known in the art. In a particularly preferred embodiment, the nucleic acid sequences of SEQ ID NO:1-1009 or novel segments or parts of the nucleic acids provided herein are used in diagnostics for identifying expressed genes or, as well known in the art and exemplified by Vollrath et al., Science 258:52-59 (1992), as expressed sequence tags for physical mapping of the human genome.

The isolated polynucleotides of the invention include, but are not limited to, a polynucleotide comprising any one of the nucleotide sequences set forth in SEQ ID NO:1-1009; a polynucleotide comprising any of the full length protein coding sequences of SEQ ID NO:1 - 1009; and a polynucleotide comprising any of the nucleotide sequences of the mature protein coding sequences of SEQ ID NO: 1- 1009. The polynucleotides of the present invention also include, but are not limited to, a polynucleotide that hybridizes under stringent hybridization conditions to (a) the complement of any one of the nucleotide sequences set forth in SEQ ID NO:1-1009; (b) a nucleotide sequence encoding any one of the amino acid sequences set forth in the Sequence Listing (*e.g.*, SEQ ID NO: 1010-2018); (c) a polynucleotide which is an allelic variant of any polynucleotides recited above; (d) a polynucleotide which encodes a species homolog (*e.g.* orthologs) of any of the proteins recited above; or (e) a polynucleotide that encodes a polypeptide comprising a specific domain or truncation of any of the polypeptides comprising an amino acid sequence set forth in the Sequence Listing.

The isolated polypeptides of the invention include, but are not limited to, a polypeptide comprising any of the amino acid sequences set forth in the Sequence Listing; or the corresponding full length or mature protein. Polypeptides of the invention also include polypeptides with biological activity that are encoded by (a) any of the polynucleotides having a nucleotide sequence set forth in SEQ ID NO:1-1009; or (b) polynucleotides that hybridize to the complement of the polynucleotides of (a) under stringent hybridization conditions. Biologically or immunologically active variants of any of the polypeptide sequences in the Sequence Listing, and "substantial equivalents" thereof (*e.g.*, with at least about 65%, 70%, 75%, 80%, 85%, 90%, 95%, 98% or 99% amino acid sequence identity) that preferably retain biological activity are also contemplated. The polypeptides of the invention may be wholly or partially chemically synthesized but are preferably produced by recombinant means using the genetically engineered cells (*e.g.* host cells) of the invention.

The invention also provides compositions comprising a polypeptide of the invention. Polypeptide compositions of the invention may further comprise an acceptable carrier, such as a hydrophilic, *e.g.*, pharmaceutically acceptable, carrier.

The invention also provides host cells transformed or transfected with a polynucleotide of the invention.

The invention also relates to methods for producing a polypeptide of the invention comprising growing a culture of the host cells of the invention in a suitable culture medium under conditions permitting expression of the desired polypeptide, and purifying the polypeptide from the culture or from the host cells. Preferred embodiments include those in which the protein produced by such process is a mature form of the protein.

Polynucleotides according to the invention have numerous applications in a variety of techniques known to those skilled in the art of molecular biology. These techniques include use as hybridization probes, use as oligomers, or primers, for PCR, use for chromosome and gene mapping, use in the recombinant production of protein, and use in generation of anti-sense DNA or RNA, their chemical analogs and the like. For example, when the expression of an mRNA is largely restricted to a particular cell or tissue type, polynucleotides of the invention can be used as hybridization probes to detect the presence of the particular cell or tissue mRNA in a sample using, *e.g., in situ* hybridization.

In other exemplary embodiments, the polynucleotides are used in diagnostics as expressed sequence tags for identifying expressed genes or, as well known in the art and exemplified by Vollrath et al., Science 258:52-59 (1992), as expressed sequence tags for physical mapping of the human genome.

The polypeptides according to the invention can be used in a variety of conventional procedures and methods that are currently applied to other proteins. For example, a polypeptide of the invention can be used to generate an antibody that specifically binds the polypeptide. Such antibodies, particularly monoclonal antibodies, are useful for detecting or quantitating the polypeptide in tissue. The polypeptides of the invention can also be used as molecular weight markers, and as a food supplement.

Methods are also provided for preventing, treating, or ameliorating a medical condition which comprises the step of administering to a mammalian subject a therapeutically effective amount of a composition comprising a polypeptide of the present invention and a pharmaceutically acceptable carrier.

In particular, the polypeptides and polynucleotides of the invention can be utilized, for example, in methods for the prevention and/or treatment of disorders involving aberrant protein expression or biological activity.

The present invention further relates to methods for detecting the presence of the polynucleotides or polypeptides of the invention in a sample. Such methods can, for example, be utilized as part of prognostic and diagnostic evaluation of disorders as recited herein and for the identification of subjects exhibiting a predisposition to such conditions. The invention provides a method for detecting the polynucleotides of the invention in a sample, comprising contacting the sample with a compound that binds to and forms a complex with the polynucleotide of interest for a period sufficient to form the complex and under conditions sufficient to form a complex and detecting the complex such that if a complex is detected, the polynucleotide of interest is detected. The invention also provides a method for detecting the polypeptides of the invention in a sample comprising contacting the sample with a compound that binds to and forms a complex with the polypeptide under conditions and for a period sufficient to form the complex and detecting the formation of the complex such that if a complex is formed, the polypeptide is detected.

The invention also provides kits comprising polynucleotide probes and/or monoclonal antibodies, and optionally quantitative standards, for carrying out methods of the invention. Furthermore, the invention provides methods for evaluating the efficacy of drugs, and monitoring the progress of patients, involved in clinical trials for the treatment of disorders as recited above.

The invention also provides methods for the identification of compounds that modulate (*i.e.*, increase or decrease) the expression or activity of the polynucleotides and/or polypeptides of the invention. Such methods can be utilized, for example, for the identification of compounds that can ameliorate symptoms of disorders as recited herein. Such methods can include, but are not limited to, assays for identifying compounds and other substances that interact with (*e.g.*, bind to) the polypeptides of the invention. The invention provides a method for identifying a compound that binds to the polypeptides of the invention comprising contacting the compound with a polypeptide of the invention in a cell for a time sufficient to form a polypeptide/compound complex, wherein the complex drives expression of a reporter gene sequence in the cell; and detecting the complex by detecting the reporter gene sequence expression such that if expression of the reporter gene is detected the compound the binds to a polypeptide of the invention is identified.

The methods of the invention also provides methods for treatment which involve the administration of the polynucleotides or polypeptides of the invention to individuals exhibiting symptoms or tendencies. In addition, the invention encompasses methods for treating diseases or disorders as recited herein comprising administering compounds and other substances that modulate the overall activity of the target gene products. Compounds and other substances can

effect such modulation either on the level of target gene/protein expression or target protein activity.

The polypeptides of the present invention and the polynucleotides encoding them are also useful for the same functions known to one of skill in the art as the polypeptides and polynucleotides to which they have homology (set forth in Table 2). If no homology is set forth for a sequence, then the polypeptides and polynucleotides of the present invention are useful for a variety of applications, as described herein, including use in arrays for detection.

## 4. DETAILED DESCRIPTION OF THE INVENTION

### 4.1 DEFINITIONS

It must be noted that as used herein and in the appended claims, the singular forms “a”, “an” and “the” include plural references unless the context clearly dictates otherwise.

The term "active" refers to those forms of the polypeptide which retain the biologic and/or immunologic activities of any naturally occurring polypeptide. According to the invention, the terms “biologically active” or “biological activity” refer to a protein or peptide having structural, regulatory or biochemical functions of a naturally occurring molecule. Likewise “immunologically active” or “immunological activity” refers to the capability of the natural, recombinant or synthetic polypeptide to induce a specific immune response in appropriate animals or cells and to bind with specific antibodies.

The term "activated cells” as used in this application are those cells which are engaged in extracellular or intracellular membrane trafficking, including the export of secretory or enzymatic molecules as part of a normal or disease process.

The terms “complementary” or “complementarity” refer to the natural binding of polynucleotides by base pairing. For example, the sequence 5’-AGT-3’ binds to the complementary sequence 3’-TCA-5’. Complementarity between two single-stranded molecules may be “partial” such that only some of the nucleic acids bind or it may be “complete” such that total complementarity exists between the single stranded molecules. The degree of complementarity between the nucleic acid strands has significant effects on the efficiency and strength of the hybridization between the nucleic acid strands.

The term “embryonic stem cells (ES)” refers to a cell that can give rise to many differentiated cell types in an embryo or an adult, including the germ cells. The term “germ line stem cells (GSCs)” refers to stem cells derived from primordial stem cells that provide a steady and continuous source of germ cells for the production of gametes. The term “primordial germ

cells (PGCs)" refers to a small population of cells set aside from other cell lineages particu... from the yolk sac, mesenteries, or gonadal ridges during embryogenesis that have the potential to differentiate into germ cells and other cells. PGCs are the source from which GSCs and ES cells are derived The PGCs, the GSCs and the ES cells are capable of self-renewal. Thus these cells not only populate the germ line and give rise to a plurality of terminally differentiated cells that comprise the adult specialized organs, but are able to regenerate themselves.

The term "expression modulating fragment," EMF, means a series of nucleotides which modulates the expression of an operably linked ORF or another EMF.

As used herein, a sequence is said to "modulate the expression of an operably linked sequence" when the expression of the sequence is altered by the presence of the EMF. EMFs include, but are not limited to, promoters, and promoter modulating sequences (inducible elements). One class of EMFs are nucleic acid fragments which induce the expression of an operably linked ORF in response to a specific regulatory factor or physiological event.

The terms "nucleotide sequence" or "nucleic acid" or "polynucleotide" or "oligonculeotide" are used interchangeably and refer to a heteropolymer of nucleotides or the sequence of these nucleotides. These phrases also refer to DNA or RNA of genomic or synthetic origin which may be single-stranded or double-stranded and may represent the sense or the antisense strand, to peptide nucleic acid (PNA) or to any DNA-like or RNA-like material. In the sequences herein A is adenine, C is cytosine, T is thymine, G is guanine and N is A, C, G or T (U). It is contemplated that where the polynucleotide is RNA, the T (thymine) in the sequences provided herein is substituted with U (uracil). Generally, nucleic acid segments provided by this invention may be assembled from fragments of the genome and short oligonucleotide linkers, or from a series of oligonucleotides, or from individual nucleotides, to provide a synthetic nucleic acid which is capable of being expressed in a recombinant transcriptional unit comprising regulatory elements derived from a microbial or viral operon, or a eukaryotic gene.

The terms "oligonucleotide fragment" or a "polynucleotide fragment", "portion," or "segment" or "probe" or "primer" are used interchangeably and refer to a sequence of nucleotide residues which are at least about 5 nucleotides, more preferably at least about 7 nucleotides, more preferably at least about 9 nucleotides, more preferably at least about 11 nucleotides and most preferably at least about 17 nucleotides. The fragment is preferably less than about 500 nucleotides, preferably less than about 200 nucleotides, more preferably less than about 100 nucleotides, more preferably less than about 50 nucleotides and most preferably less than 30 nucleotides. Preferably the probe is from about 6 nucleotides to about 200 nucleotides, preferably from about 15 to about 50 nucleotides, more preferably from about 17 to 30 nucleotides and most preferably from about 20 to 25 nucleotides. Preferably the fragments can

be used in polymerase chain reaction (PCR), various hybridization procedures or microarray procedures to identify or amplify identical or related parts of mRNA or DNA molecules. A fragment or segment may uniquely identify each polynucleotide sequence of the present invention. Preferably the fragment comprises a sequence substantially similar to any one of SEQ ID NOs:1-1009.

Probes may, for example, be used to determine whether specific mRNA molecules are present in a cell or tissue or to isolate similar nucleic acid sequences from chromosomal DNA as described by Walsh et al. (Walsh, P.S. et al., 1992, PCR Methods Appl 1:241-250). They may be labeled by nick translation, Klenow fill-in reaction, PCR, or other methods well known in the art. Probes of the present invention, their preparation and/or labeling are elaborated in Sambrook, J. et al., 1989, Molecular Cloning: A Laboratory Manual, Cold Spring Harbor Laboratory, NY; or Ausubel, F.M. et al., 1989, Current Protocols in Molecular Biology, John Wiley & Sons, New York NY, both of which are incorporated herein by reference in their entirety.

The nucleic acid sequences of the present invention also include the sequence information from the nucleic acid sequences of SEQ ID NO:1-1009. The sequence information can be a segment of any one of SEQ ID NO:1-1009 that uniquely identifies or represents the sequence information of that sequence of SEQ ID NO:1-1009. One such segment can be a twenty-mer nucleic acid sequence because the probability that a twenty-mer is fully matched in the human genome is 1 in 300. In the human genome, there are three billion base pairs in one set of chromosomes. Because $4^{20}$ possible twenty-mers exist, there are 300 times more twenty-mers than there are base pairs in a set of human chromosomes. Using the same analysis, the probability for a seventeen-mer to be fully matched in the human genome is approximately 1 in 5. When these segments are used in arrays for expression studies, fifteen-mer segments can be used. The probability that the fifteen-mer is fully matched in the expressed sequences is also approximately one in five because expressed sequences comprise less than approximately 5% of the entire genome sequence.

Similarly, when using sequence information for detecting a single mismatch, a segment can be a twenty-five mer. The probability that the twenty-five mer would appear in a human genome with a single mismatch is calculated by multiplying the probability for a full match ($1 \div 4^{25}$) times the increased probability for mismatch at each nucleotide position (3 x 25). The probability that an eighteen mer with a single mismatch can be detected in an array for expression studies is approximately one in five. The probability that a twenty-mer with a single mismatch can be detected in a human genome is approximately one in five.

The term "open reading frame," ORF, means a series of nucleotide triplets coding for amino acids without any termination codons and is a sequence translatable into protein.

The terms “operably linked” or “operably associated” refer to functionally related nucleic acid sequences. For example, a promoter is operably associated or operably linked with a coding sequence if the promoter controls the transcription of the coding sequence. While operably linked nucleic acid sequences can be contiguous and in the same reading frame, certain genetic elements *e.g.* repressor genes are not contiguously linked to the coding sequence but still control transcription/translation of the coding sequence.

The term “pluripotent” refers to the capability of a cell to differentiate into a number of differentiated cell types that are present in an adult organism. A pluripotent cell is restricted in its differentiation capability in comparison to a totipotent cell.

The terms “polypeptide” or “peptide” or “amino acid sequence” refer to an oligopeptide, peptide, polypeptide or protein sequence or fragment thereof and to naturally occurring or synthetic molecules. A polypeptide "fragment," "portion," or "segment" is a stretch of amino acid residues of at least about 5 amino acids, preferably at least about 7 amino acids, more preferably at least about 9 amino acids and most preferably at least about 17 or more amino acids. The peptide preferably is not greater than about 200 amino acids, more preferably less than 150 amino acids and most preferably less than 100 amino acids. Preferably the peptide is from about 5 to about 200 amino acids. To be active, any polypeptide must have sufficient length to display biological and/or immunological activity.

The term "naturally occurring polypeptide" refers to polypeptides produced by cells that have not been genetically engineered and specifically contemplates various polypeptides arising from post-translational modifications of the polypeptide including, but not limited to, acetylation, carboxylation, glycosylation, phosphorylation, lipidation and acylation.

The term “translated protein coding portion” means a sequence which encodes for the full length protein which may include any leader sequence or any processing sequence.

The term “mature protein coding sequence” means a sequence which encodes a peptide or protein without a signal or leader sequence. The “mature protein portion” means that portion of the protein which does not include a signal or leader sequence. The peptide may have been produced by processing in the cell which removes any leader/signal sequence. The mature protein portion may or may not include the initial methionine residue. The methionine residue may be removed from the protein during processing in the cell. The peptide may be produced synthetically or the protein may have been produced using a polynucleotide only encoding for the mature protein coding sequence.

The term "derivative" refers to polypeptides chemically modified by such techniques as ubiquitination, labeling (*e.g.*, with radionuclides or various enzymes), covalent polymer attachment such as pegylation (derivatization with polyethylene glycol) and insertion or substitution by chemical synthesis of amino acids such as ornithine, which do not normally occur in human proteins.

The term "variant"(or "analog") refers to any polypeptide differing from naturally occurring polypeptides by amino acid insertions, deletions, and substitutions, created using, *e g.*, recombinant DNA techniques. Guidance in determining which amino acid residues may be replaced, added or deleted without abolishing activities of interest, may be found by comparing the sequence of the particular polypeptide with that of homologous peptides and minimizing the number of amino acid sequence changes made in regions of high homology (conserved regions) or by replacing amino acids with consensus sequence.

Alternatively, recombinant variants encoding these same or similar polypeptides may be synthesized or selected by making use of the "redundancy" in the genetic code. Various codon substitutions, such as the silent changes which produce various restriction sites, may be introduced to optimize cloning into a plasmid or viral vector or expression in a particular prokaryotic or eukaryotic system. Mutations in the polynucleotide sequence may be reflected in the polypeptide or domains of other peptides added to the polypeptide to modify the properties of any part of the polypeptide, to change characteristics such as ligand-binding affinities, interchain affinities, or degradation/turnover rate.

Preferably, amino acid "substitutions" are the result of replacing one amino acid with another amino acid having similar structural and/or chemical properties, *i.e.*, conservative amino acid replacements. "Conservative" amino acid substitutions may be made on the basis of similarity in polarity, charge, solubility, hydrophobicity, hydrophilicity, and/or the amphipathic nature of the residues involved. For example, nonpolar (hydrophobic) amino acids include alanine, leucine, isoleucine, valine, proline, phenylalanine, tryptophan, and methionine; polar neutral amino acids include glycine, serine, threonine, cysteine, tyrosine, asparagine, and glutamine; positively charged (basic) amino acids include arginine, lysine, and histidine; and negatively charged (acidic) amino acids include aspartic acid and glutamic acid. "Insertions" or "deletions" are preferably in the range of about 1 to 20 amino acids, more preferably 1 to 10 amino acids. The variation allowed may be experimentally determined by systematically making insertions, deletions, or substitutions of amino acids in a polypeptide molecule using recombinant DNA techniques and assaying the resulting recombinant variants for activity.

Alternatively, where alteration of function is desired, insertions, deletions or non-conservative alterations can be engineered to produce altered polypeptides. Such alterations

can, for example, alter one or more of the biological functions or biochemical characteristics of the polypeptides of the invention. For example, such alterations may change polypeptide characteristics such as ligand-binding affinities, interchain affinities, or degradation/turnover rate. Further, such alterations can be selected so as to generate polypeptides that are better suited for expression, scale up and the like in the host cells chosen for expression. For example, cysteine residues can be deleted or substituted with another amino acid residue in order to eliminate disulfide bridges.

The terms "purified" or “substantially purified” as used herein denotes that the indicated nucleic acid or polypeptide is present in the substantial absence of other biological macromolecules, *e.g.*, polynucleotides, proteins, and the like. In one embodiment, the polynucleotide or polypeptide is purified such that it constitutes at least 95% by weight, more preferably at least 99% by weight, of the indicated biological macromolecules present (but water, buffers, and other small molecules, especially molecules having a molecular weight of less than 1000 daltons, can be present).

The term "isolated" as used herein refers to a nucleic acid or polypeptide separated from at least one other component (*e.g.*, nucleic acid or polypeptide) present with the nucleic acid or polypeptide in its natural source. In one embodiment, the nucleic acid or polypeptide is found in the presence of (if anything) only a solvent, buffer, ion, or other component normally present in a solution of the same. The terms "isolated" and "purified" do not encompass nucleic acids or polypeptides present in their natural source.

The term "recombinant," when used herein to refer to a polypeptide or protein, means that a polypeptide or protein is derived from recombinant (*e.g.*, microbial, insect, or mammalian) expression systems. "Microbial" refers to recombinant polypeptides or proteins made in bacterial or fungal (*e.g.*, yeast) expression systems. As a product, "recombinant microbial" defines a polypeptide or protein essentially free of native endogenous substances and unaccompanied by associated native glycosylation. Polypeptides or proteins expressed in most bacterial cultures, *e.g., E. coli*, will be free of glycosylation modifications; polypeptides or proteins expressed in yeast will have a glycosylation pattern in general different from those expressed in mammalian cells.

The term "recombinant expression vehicle or vector" refers to a plasmid or phage or virus or vector, for expressing a polypeptide from a DNA (RNA) sequence. An expression vehicle can comprise a transcriptional unit comprising an assembly of (1) a genetic element or elements having a regulatory role in gene expression, for example, promoters or enhancers, (2) a structural or coding sequence which is transcribed into mRNA and translated into protein, and (3) appropriate transcription initiation and termination sequences. Structural units intended for use

in yeast or eukaryotic expression systems preferably include a leader sequence enabling extracellular secretion of translated protein by a host cell. Alternatively, where recombinant protein is expressed without a leader or transport sequence, it may include an amino terminal methionine residue. This residue may or may not be subsequently cleaved from the expressed recombinant protein to provide a final product.

The term "recombinant expression system" means host cells which have stably integrated a recombinant transcriptional unit into chromosomal DNA or carry the recombinant transcriptional unit extrachromosomally. Recombinant expression systems as defined herein will express heterologous polypeptides or proteins upon induction of the regulatory elements linked to the DNA segment or synthetic gene to be expressed. This term also means host cells which have stably integrated a recombinant genetic element or elements having a regulatory role in gene expression, for example, promoters or enhancers. Recombinant expression systems as defined herein will express polypeptides or proteins endogenous to the cell upon induction of the regulatory elements linked to the endogenous DNA segment or gene to be expressed. The cells can be prokaryotic or eukaryotic.

The term "secreted" includes a protein that is transported across or through a membrane, including transport as a result of signal sequences in its amino acid sequence when it is expressed in a suitable host cell. "Secreted" proteins include without limitation proteins secreted wholly (*e.g.*, soluble proteins) or partially (*e.g.*, receptors) from the cell in which they are expressed. "Secreted" proteins also include without limitation proteins that are transported across the membrane of the endoplasmic reticulum. "Secreted" proteins are also intended to include proteins containing non-typical signal sequences (*e.g.* Interleukin-1 Beta, see Krasney, P.A. and Young, P.R. (1992) Cytokine 4(2):134 -143) and factors released from damaged cells (*e.g.* Interleukin-1 Receptor Antagonist, see Arend, W.P. et. al. (1998) Annu. Rev. Immunol. 16:27-55)

Where desired, an expression vector may be designed to contain a "signal or leader sequence" which will direct the polypeptide through the membrane of a cell. Such a sequence may be naturally present on the polypeptides of the present invention or provided from heterologous protein sources by recombinant DNA techniques.

The term "stringent" is used to refer to conditions that are commonly understood in the art as stringent. Stringent conditions can include highly stringent conditions (*i.e.*, hybridization to filter-bound DNA in 0.5 M $NaHPO_4$, 7% sodium dodecyl sulfate (SDS), 1 mM EDTA at 65°C, and washing in 0.1X SSC/0.1% SDS at 68°C), and moderately stringent conditions (*i.e.*, washing in 0.2X SSC/0.1% SDS at 42°C). Other exemplary hybridization conditions are described herein in the examples.

In instances of hybridization of deoxyoligonucleotides, additional exemplary stringent hybridization conditions include washing in 6X SSC/0.05% sodium pyrophosphate at 37°C (for 14-base oligonucleotides), 48°C (for 17-base oligos), 55°C (for 20-base oligonucleotides), and 60°C (for 23-base oligonucleotides).

As used herein, "substantially equivalent" can refer both to nucleotide and amino acid sequences, for example a mutant sequence, that varies from a reference sequence by one or more substitutions, deletions, or additions, the net effect of which does not result in an adverse functional dissimilarity between the reference and subject sequences. Typically, such a substantially equivalent sequence varies from one of those listed herein by no more than about 35% (*i.e.*, the number of individual residue substitutions, additions, and/or deletions in a substantially equivalent sequence, as compared to the corresponding reference sequence, divided by the total number of residues in the substantially equivalent sequence is about 0.35 or less). Such a sequence is said to have 65% sequence identity to the listed sequence. In one embodiment, a substantially equivalent, *e.g.*, mutant, sequence of the invention varies from a listed sequence by no more than 30% (70% sequence identity); in a variation of this embodiment, by no more than 25% (75% sequence identity); and in a further variation of this embodiment, by no more than 20% (80% sequence identity) and in a further variation of this embodiment, by no more than 10% (90% sequence identity) and in a further variation of this embodiment, by no more that 5% (95% sequence identity). Substantially equivalent, *e.g.*, mutant, amino acid sequences according to the invention preferably have at least 80% sequence identity with a listed amino acid sequence, more preferably at least 85% sequence identity, more preferably at least 90% sequence identity, more preferably at least 95% identity, more preferably at least 98% identity, and most preferably at least 99% identity. Substantially equivalent nucleotide sequences of the invention can have lower percent sequence identities, taking into account, for example, the redundancy or degeneracy of the genetic code. Preferably, nucleotide sequence has at least about 65% identity, more preferably at least about 75% identity, more preferably at least about 80% sequence identity, more preferably at least about 85% sequence identity, more preferably at least about 90% sequence identity, and most preferably at least about 95% identity, more preferably at least about 98% sequence identity, and most preferably at least about 99% sequence identity. For the purposes of the present invention, sequences having substantially equivalent biological activity and substantially equivalent expression characteristics are considered substantially equivalent. For the purposes of determining equivalence, truncation of the mature sequence (*e.g.*, via a mutation which creates a spurious stop codon) should be disregarded. Sequence identity may be determined, *e.g.*, using the Jotun Hein method (Hein, J.

(1990) Methods Enzymol. 183:626-645). Identity between sequences can also be determined by other methods known in the art, *e.g.* by varying hybridization conditions.

The term "totipotent" refers to the capability of a cell to differentiate into all of the cell types of an adult organism.

The term "transformation" means introducing DNA into a suitable host cell so that the DNA is replicable, either as an extrachromosomal element, or by chromosomal integration. The term "transfection" refers to the taking up of an expression vector by a suitable host cell, whether or not any coding sequences are in fact expressed. The term "infection" refers to the introduction of nucleic acids into a suitable host cell by use of a virus or viral vector.

As used herein, an "uptake modulating fragment," UMF, means a series of nucleotides which mediate the uptake of a linked DNA fragment into a cell. UMFs can be readily identified using known UMFs as a target sequence or target motif with the computer-based systems described below. The presence and activity of a UMF can be confirmed by attaching the suspected UMF to a marker sequence. The resulting nucleic acid molecule is then incubated with an appropriate host under appropriate conditions and the uptake of the marker sequence is determined. As described above, a UMF will increase the frequency of uptake of a linked marker sequence.

Each of the above terms is meant to encompass all that is described for each, unless the context dictates otherwise.

### 4.2 NUCLEIC ACIDS OF THE INVENTION

Nucleotide sequences of the invention are set forth in the Sequence Listing.

The isolated polynucleotides of the invention include a polynucleotide comprising the nucleotide sequences of SEQ ID NO:1-1009 ; a polynucleotide encoding any one of the peptide sequences of SEQ ID NO:1010-2018; and a polynucleotide comprising the nucleotide sequence encoding the mature protein coding sequence of the polypeptides of any one of SEQ ID NO:1010-2018. The polynucleotides of the present invention also include, but are not limited to, a polynucleotide that hybridizes under stringent conditions to (a) the complement of any of the nucleotides sequences of SEQ ID NO:1-1009 ; (b) nucleotide sequences encoding any one of the amino acid sequences set forth in the Sequence Listing; (c) a polynucleotide which is an allelic variant of any polynucleotide recited above; (d) a polynucleotide which encodes a species homolog of any of the proteins recited above; or (e) a polynucleotide that encodes a polypeptide comprising a specific domain or truncation of the polypeptides of SEQ ID NO: 1010-2018. Domains of interest may depend on the nature of the encoded polypeptide; *e.g.*, domains in receptor-like polypeptides include ligand-binding, extracellular, transmembrane, or cytoplasmic

domains, or combinations thereof; domains in immunoglobulin-like proteins include the variable immunoglobulin-like domains; domains in enzyme-like polypeptides include catalytic and substrate binding domains; and domains in ligand polypeptides include receptor-binding domains.

The polynucleotides of the invention include naturally occurring or wholly or partially synthetic DNA, *e.g.*, cDNA and genomic DNA, and RNA, *e.g.*, mRNA. The polynucleotides may include all of the coding region of the cDNA or may represent a portion of the coding region of the cDNA.

The present invention also provides genes corresponding to the cDNA sequences disclosed herein. The corresponding genes can be isolated in accordance with known methods using the sequence information disclosed herein. Such methods include the preparation of probes or primers from the disclosed sequence information for identification and/or amplification of genes in appropriate genomic libraries or other sources of genomic materials. Further 5' and 3' sequence can be obtained using methods known in the art. For example, full length cDNA or genomic DNA that corresponds to any of the polynucleotides of SEQ ID NO:1-1009 can be obtained by screening appropriate cDNA or genomic DNA libraries under suitable hybridization conditions using any of the polynucleotides of SEQ ID NO:1-1009 or a portion thereof as a probe. Alternatively, the polynucleotides of SEQ ID NO:1-1009 may be used as the basis for suitable primer(s) that allow identification and/or amplification of genes in appropriate genomic DNA or cDNA libraries.

The nucleic acid sequences of the invention can be assembled from ESTs and sequences (including cDNA and genomic sequences) obtained from one or more public databases, such as dbEST, gbpri, and UniGene. The EST sequences can provide identifying sequence information, representative fragment or segment information, or novel segment information for the full-length gene.

The polynucleotides of the invention also provide polynucleotides including nucleotide sequences that are substantially equivalent to the polynucleotides recited above. Polynucleotides according to the invention can have, *e.g.*, at least about 65%, at least about 70%, at least about 75%, at least about 80%, 81%, 82%, 83%, 84%, more typically at least about 85%, 86%, 87%, 88%, 89%, more typically at least about 90%, 91%, 92%, 93%, 94%, and even more typically at least about 95%, 96%, 97%, 98%, 99%, sequence identity to a polynucleotide recited above.

Included within the scope of the nucleic acid sequences of the invention are nucleic acid sequence fragments that hybridize under stringent conditions to any of the nucleotide sequences of SEQ ID NO:1-1009, or complements thereof, which fragment is greater than about 5 nucleotides, preferably 7 nucleotides, more preferably greater than 9 nucleotides and most preferably greater than 17 nucleotides. Fragments of, *e.g.* 15, 17, or 20 nucleotides or more that

are selective for (*i.e.* specifically hybridize to any one of the polynucleotides of the invention) are contemplated. Probes capable of specifically hybridizing to a polynucleotide can differentiate polynucleotide sequences of the invention from other polynucleotide sequences in the same family of genes or can differentiate human genes from genes of other species, and are preferably based on unique nucleotide sequences.

The sequences falling within the scope of the present invention are not limited to these specific sequences, but also include allelic and species variations thereof. Allelic and species variations can be routinely determined by comparing the sequence provided SEQ ID NO:1-1009, a representative fragment thereof, or a nucleotide sequence at least 90% identical, preferably 95% identical, to SEQ ID NO:1-1009 with a sequence from another isolate of the same species. Furthermore, to accommodate codon variability, the invention includes nucleic acid molecules coding for the same amino acid sequences as do the specific ORFs disclosed herein. In other words, in the coding region of an ORF, substitution of one codon for another codon that encodes the same amino acid is expressly contemplated.

The nearest neighbor or homology result for the nucleic acids of the present invention, including SEQ ID NO:1-1009, can be obtained by searching a database using an algorithm or a program. Preferably, a BLAST which stands for Basic Local Alignment Search Tool is used to search for local sequence alignments (Altshul, S.F. J Mol. Evol. 36 290-300 (1993) and Altschul S.F. et al. J. Mol. Biol. 21:403-410 (1990)). Alternatively a FASTA version 3 search against Genpept, using Fastxy algorithm.

Species homologs (or orthologs) of the disclosed polynucleotides and proteins are also provided by the present invention. Species homologs may be isolated and identified by making suitable probes or primers from the sequences provided herein and screening a suitable nucleic acid source from the desired species.

The invention also encompasses allelic variants of the disclosed polynucleotides or proteins; that is, naturally-occurring alternative forms of the isolated polynucleotide which also encode proteins which are identical, homologous or related to that encoded by the polynucleotides.

The nucleic acid sequences of the invention are further directed to sequences which encode variants of the described nucleic acids. These amino acid sequence variants may be prepared by methods known in the art by introducing appropriate nucleotide changes into a native or variant polynucleotide. There are two variables in the construction of amino acid sequence variants: the location of the mutation and the nature of the mutation. Nucleic acids encoding the amino acid sequence variants are preferably constructed by mutating the polynucleotide to encode an amino acid sequence that does not occur in nature. These nucleic

acid alterations can be made at sites that differ in the nucleic acids from different species (variable positions) or in highly conserved regions (constant regions). Sites at such locations will typically be modified in series, *e.g.*, by substituting first with conservative choices (*e.g.*, hydrophobic amino acid to a different hydrophobic amino acid) and then with more distant choices (*e.g.*, hydrophobic amino acid to a charged amino acid), and then deletions or insertions may be made at the target site. Amino acid sequence deletions generally range from about 1 to 30 residues, preferably about 1 to 10 residues, and are typically contiguous. Amino acid insertions include amino- and/or carboxyl-terminal fusions ranging in length from one to one hundred or more residues, as well as intrasequence insertions of single or multiple amino acid residues. Intrasequence insertions may range generally from about 1 to 10 amino residues, preferably from 1 to 5 residues. Examples of terminal insertions include the heterologous signal sequences necessary for secretion or for intracellular targeting in different host cells and sequences such as FLAG or poly-histidine sequences useful for purifying the expressed protein.

In a preferred method, polynucleotides encoding the novel amino acid sequences are changed via site-directed mutagenesis. This method uses oligonucleotide sequences to alter a polynucleotide to encode the desired amino acid variant, as well as sufficient adjacent nucleotides on both sides of the changed amino acid to form a stable duplex on either side of the site of being changed. In general, the techniques of site-directed mutagenesis are well known to those of skill in the art and this technique is exemplified by publications such as, Edelman et al., *DNA* 2:183 (1983). A versatile and efficient method for producing site-specific changes in a polynucleotide sequence was published by Zoller and Smith, *Nucleic Acids Res.* 10:6487-6500 (1982). PCR may also be used to create amino acid sequence variants of the novel nucleic acids. When small amounts of template DNA are used as starting material, primer(s) that differs slightly in sequence from the corresponding region in the template DNA can generate the desired amino acid variant. PCR amplification results in a population of product DNA fragments that differ from the polynucleotide template encoding the polypeptide at the position specified by the primer. The product DNA fragments replace the corresponding region in the plasmid and this gives a polynucleotide encoding the desired amino acid variant.

A further technique for generating amino acid variants is the cassette mutagenesis technique described in Wells et al., *Gene* 34:315 (1985); and other mutagenesis techniques well known in the art, such as, for example, the techniques in Sambrook et al., supra, and *Current Protocols in Molecular Biology*, Ausubel et al. Due to the inherent degeneracy of the genetic code, other DNA sequences which encode substantially the same or a functionally equivalent amino acid sequence may be used in the practice of the invention for the cloning and expression

of these novel nucleic acids. Such DNA sequences include those which are capable of hybridizing to the appropriate novel nucleic acid sequence under stringent conditions.

Polynucleotides encoding preferred polypeptide truncations of the invention can be used to generate polynucleotides encoding chimeric or fusion proteins comprising one or more domains of the invention and heterologous protein sequences.

The polynucleotides of the invention additionally include the complement of any of the polynucleotides recited above. The polynucleotide can be DNA (genomic, cDNA, amplified, or synthetic) or RNA. Methods and algorithms for obtaining such polynucleotides are well known to those of skill in the art and can include, for example, methods for determining hybridization conditions that can routinely isolate polynucleotides of the desired sequence identities.

In accordance with the invention, polynucleotide sequences comprising the mature protein coding sequences corresponding to any one of SEQ ID NO:1-1009, or functional equivalents thereof, may be used to generate recombinant DNA molecules that direct the expression of that nucleic acid, or a functional equivalent thereof, in appropriate host cells. Also included are the cDNA inserts of any of the clones identified herein.

A polynucleotide according to the invention can be joined to any of a variety of other nucleotide sequences by well-established recombinant DNA techniques (see Sambrook J et al. (1989) Molecular Cloning: A Laboratory Manual, Cold Spring Harbor Laboratory, NY). Useful nucleotide sequences for joining to polynucleotides include an assortment of vectors, *e.g.*, plasmids, cosmids, lambda phage derivatives, phagemids, and the like, that are well known in the art. Accordingly, the invention also provides a vector including a polynucleotide of the invention and a host cell containing the polynucleotide. In general, the vector contains an origin of replication functional in at least one organism, convenient restriction endonuclease sites, and a selectable marker for the host cell. Vectors according to the invention include expression vectors, replication vectors, probe generation vectors, and sequencing vectors. A host cell according to the invention can be a prokaryotic or eukaryotic cell and can be a unicellular organism or part of a multicellular organism.

The present invention further provides recombinant constructs comprising a nucleic acid having any of the nucleotide sequences of SEQ ID NO:1-1009 or a fragment thereof or any other polynucleotides of the invention. In one embodiment, the recombinant constructs of the present invention comprise a vector, such as a plasmid or viral vector, into which a nucleic acid having any of the nucleotide sequences of SEQ ID NO:1-1009 or a fragment thereof is inserted, in a forward or reverse orientation. In the case of a vector comprising one of the ORFs of the present invention, the vector may further comprise regulatory sequences, including for example, a promoter, operably linked to the ORF. Large numbers of suitable vectors and promoters are

known to those of skill in the art and are commercially available for generating the recombinant constructs of the present invention. The following vectors are provided by way of example. Bacterial: pBs, phagescript, PsiX174, pBluescript SK, pBs KS, pNH8a, pNH16a, pNH18a, pNH46a (Stratagene); pTrc99A, pKK223-3, pKK233-3, pDR540, pRIT5 (Pharmacia). Eukaryotic: pWLneo, pSV2cat, pOG44, PXTI, pSG (Stratagene) pSVK3, pBPV, pMSG, pSVL (Pharmacia).

The isolated polynucleotide of the invention may be operably linked to an expression control sequence such as the pMT2 or pED expression vectors disclosed in Kaufman et al., *Nucleic Acids Res.* 19, 4485-4490 (1991), in order to produce the protein recombinantly. Many suitable expression control sequences are known in the art. General methods of expressing recombinant proteins are also known and are exemplified in R. Kaufman, *Methods in Enzymology* 185, 537-566 (1990). As defined herein "operably linked" means that the isolated polynucleotide of the invention and an expression control sequence are situated within a vector or cell in such a way that the protein is expressed by a host cell which has been transformed (transfected) with the ligated polynucleotide/expression control sequence.

Promoter regions can be selected from any desired gene using CAT (chloramphenicol transferase) vectors or other vectors with selectable markers. Two appropriate vectors are pKK232-8 and pCM7. Particular named bacterial promoters include lacI, lacZ, T3, T7, gpt, lambda PR, and trc. Eukaryotic promoters include CMV immediate early, HSV thymidine kinase, early and late SV40, LTRs from retrovirus, and mouse metallothionein-I. Selection of the appropriate vector and promoter is well within the level of ordinary skill in the art. Generally, recombinant expression vectors will include origins of replication and selectable markers permitting transformation of the host cell, *e.g.*, the ampicillin resistance gene of *E. coli* and *S. cerevisiae* TRP1 gene, and a promoter derived from a highly-expressed gene to direct transcription of a downstream structural sequence. Such promoters can be derived from operons encoding glycolytic enzymes such as 3-phosphoglycerate kinase (PGK), a-factor, acid phosphatase, or heat shock proteins, among others. The heterologous structural sequence is assembled in appropriate phase with translation initiation and termination sequences, and preferably, a leader sequence capable of directing secretion of translated protein into the periplasmic space or extracellular medium. Optionally, the heterologous sequence can encode a fusion protein including an amino terminal identification peptide imparting desired characteristics, *e.g.*, stabilization or simplified purification of expressed recombinant product. Useful expression vectors for bacterial use are constructed by inserting a structural DNA sequence encoding a desired protein together with suitable translation initiation and termination signals in operable reading phase with a functional promoter. The vector will comprise one or

more phenotypic selectable markers and an origin of replication to ensure maintenance of the vector and to, if desirable, provide amplification within the host. Suitable prokaryotic hosts for transformation include *E. coli*, *Bacillus subtilis*, *Salmonella typhimurium* and various species within the genera *Pseudomonas*, *Streptomyces*, and *Staphylococcus*, although others may also be employed as a matter of choice.

As a representative but non-limiting example, useful expression vectors for bacterial use can comprise a selectable marker and bacterial origin of replication derived from commercially available plasmids comprising genetic elements of the well known cloning vector pBR322 (ATCC 37017). Such commercial vectors include, for example, pKK223-3 (Pharmacia Fine Chemicals, Uppsala, Sweden) and GEM 1 (Promega Biotech, Madison, WI, USA). These pBR322 "backbone" sections are combined with an appropriate promoter and the structural sequence to be expressed. Following transformation of a suitable host strain and growth of the host strain to an appropriate cell density, the selected promoter is induced or derepressed by appropriate means (*e.g.*, temperature shift or chemical induction) and cells are cultured for an additional period. Cells are typically harvested by centrifugation, disrupted by physical or chemical means, and the resulting crude extract retained for further purification.

Polynucleotides of the invention can also be used to induce immune responses. For example, as described in Fan et al., *Nat. Biotech.* 17:870-872 (1999), incorporated herein by reference, nucleic acid sequences encoding a polypeptide may be used to generate antibodies against the encoded polypeptide following topical administration of naked plasmid DNA or following injection, and preferably intramuscular injection of the DNA. The nucleic acid sequences are preferably inserted in a recombinant expression vector and may be in the form of naked DNA.

### 4.3 ANTISENSE

Another aspect of the invention pertains to isolated antisense nucleic acid molecules that are hybridizable to or complementary to the nucleic acid molecule comprising the nucleotide sequence of SEQ ID NO:1-1009, or fragments, analogs or derivatives thereof. An "antisense" nucleic acid comprises a nucleotide sequence that is complementary to a "sense" nucleic acid encoding a protein, *e.g.*, complementary to the coding strand of a double-stranded cDNA molecule or complementary to an mRNA sequence. In specific aspects, antisense nucleic acid molecules are provided that comprise a sequence complementary to at least about 10, 25, 50, 100, 250 or 500 nucleotides or an entire coding strand, or to only a portion thereof. Nucleic acid molecules encoding fragments, homologs, derivatives and analogs of a protein of any of SEQ ID

NO:1010-2018 or antisense nucleic acids complementary to a nucleic acid sequence of SEQ ID NO:1-1009 are additionally provided.

In one embodiment, an antisense nucleic acid molecule is antisense to a "coding region" of the coding strand of a nucleotide sequence of the invention. The term "coding region" refers to the region of the nucleotide sequence comprising codons which are translated into amino acid residues. In another embodiment, the antisense nucleic acid molecule is antisense to a "noncoding region" of the coding strand of a nucleotide sequence of the invention. The term "noncoding region" refers to 5' and 3' sequences which flank the coding region that are not translated into amino acids (*i.e.*, also referred to as 5' and 3' untranslated regions).

Given the coding strand sequences encoding a nucleic acid disclosed herein (*e.g.*, SEQ ID NO:1-1009), antisense nucleic acids of the invention can be designed according to the rules of Watson and Crick or Hoogsteen base pairing. The antisense nucleic acid molecule can be complementary to the entire coding region of a mRNA, but more preferably is an oligonucleotide that is antisense to only a portion of the coding or noncoding region of a mRNA. For example, the antisense oligonucleotide can be complementary to the region surrounding the translation start site of a mRNA. An antisense oligonucleotide can be, for example, about 5, 10, 15, 20, 25, 30, 35, 40, 45 or 50 nucleotides in length. An antisense nucleic acid of the invention can be constructed using chemical synthesis or enzymatic ligation reactions using procedures known in the art. For example, an antisense nucleic acid (*e.g.*, an antisense oligonucleotide) can be chemically synthesized using naturally occurring nucleotides or variously modified nucleotides designed to increase the biological stability of the molecules or to increase the physical stability of the duplex formed between the antisense and sense nucleic acids, *e.g.*, phosphorothioate derivatives and acridine substituted nucleotides can be used.

Examples of modified nucleotides that can be used to generate the antisense nucleic acid include: 5-fluorouracil, 5-bromouracil, 5-chlorouracil, 5-iodouracil, hypoxanthine, xanthine, 4-acetylcytosine, 5-(carboxyhydroxylmethyl) uracil, 5-carboxymethylaminomethyl-2-thiouridine, 5-carboxymethylaminomethyluracil, dihydrouracil, beta-D-galactosylqueosine, inosine, N6-isopentenyladenine, 1-methylguanine, 1-methylinosine, 2,2-dimethylguanine, 2-methyladenine, 2-methylguanine, 3-methylcytosine, 5-methylcytosine, N6-adenine, 7-methylguanine, 5-methylaminomethyluracil, 5-methoxyaminomethyl-2-thiouracil, beta-D-mannosylqueosine, 5'-methoxycarboxymethyluracil, 5-methoxyuracil, 2-methylthio-N6-isopentenyladenine, uracil-5-oxyacetic acid (v), wybutoxosine, pseudouracil, queosine, 2-thiocytosine, 5-methyl-2-thiouracil, 2-thiouracil, 4-thiouracil, 5-methyluracil, uracil-5-oxyacetic acid methylester, uracil-5-oxyacetic acid (v), 5-methyl-2-thiouracil, 3-(3-amino-3-N-2-carboxypropyl) uracil, (acp3)w, and 2,6-diaminopurine. Alternatively, the

antisense nucleic acid can be produced biologically using an expression vector into which a nucleic acid has been subcloned in an antisense orientation (*i.e.*, RNA transcribed from the inserted nucleic acid will be of an antisense orientation to a target nucleic acid of interest, described further in the following subsection).

The antisense nucleic acid molecules of the invention are typically administered to a subject or generated *in situ* such that they hybridize with or bind to cellular mRNA and/or genomic DNA encoding a protein according to the invention to thereby inhibit expression of the protein, *e.g.*, by inhibiting transcription and/or translation. The hybridization can be by conventional nucleotide complementarity to form a stable duplex, or, for example, in the case of an antisense nucleic acid molecule that binds to DNA duplexes, through specific interactions in the major groove of the double helix. An example of a route of administration of antisense nucleic acid molecules of the invention includes direct injection at a tissue site. Alternatively, antisense nucleic acid molecules can be modified to target selected cells and then administered systemically. For example, for systemic administration, antisense molecules can be modified such that they specifically bind to receptors or antigens expressed on a selected cell surface, *e.g.*, by linking the antisense nucleic acid molecules to peptides or antibodies that bind to cell surface receptors or antigens. The antisense nucleic acid molecules can also be delivered to cells using the vectors described herein. To achieve sufficient intracellular concentrations of antisense molecules, vector constructs in which the antisense nucleic acid molecule is placed under the control of a strong pol II or pol III promoter are preferred.

In yet another embodiment, the antisense nucleic acid molecule of the invention is an α-anomeric nucleic acid molecule. An α-anomeric nucleic acid molecule forms specific double-stranded hybrids with complementary RNA in which, contrary to the usual β-units, the strands run parallel to each other (Gaultier *et al.* (1987) *Nucleic Acids Res* 15: 6625-6641). The antisense nucleic acid molecule can also comprise a 2'-o-methylribonucleotide (Inoue *et al.* (1987) *Nucleic Acids Res* 15: 6131-6148) or a chimeric RNA -DNA analogue (Inoue *et al.* (1987) *FEBS Lett* 215: 327-330).

### 4.4 RIBOZYMES AND PNA MOIETIES

In still another embodiment, an antisense nucleic acid of the invention is a ribozyme. Ribozymes are catalytic RNA molecules with ribonuclease activity that are capable of cleaving a single-stranded nucleic acid, such as a mRNA, to which they have a complementary region. Thus, ribozymes (*e.g.*, hammerhead ribozymes (described in Haselhoff and Gerlach (1988) *Nature* 334:585-591)) can be used to catalytically cleave a mRNA transcripts to thereby inhibit translation of a mRNA. A ribozyme having specificity for a nucleic acid of the invention can be

designed based upon the nucleotide sequence of a DNA disclosed herein (*i.e.*, SEQ ID NO:1-1009). For example, a derivative of a Tetrahymena L-19 IVS RNA can be constructed in which the nucleotide sequence of the active site is complementary to the nucleotide sequence to be cleaved in a SECX-encoding mRNA. See, *e.g.*, Cech *et al.* U.S. Pat. No. 4,987,071; and Cech *et al.* U.S. Pat. No. 5,116,742. Alternatively, SECX mRNA can be used to select a catalytic RNA having a specific ribonuclease activity from a pool of RNA molecules. See, *e.g.*, Bartel *et al.*, (1993) *Science* 261:1411-1418.

Alternatively, gene expression can be inhibited by targeting nucleotide sequences complementary to the regulatory region (*e.g.*, promoter and/or enhancers) to form triple helical structures that prevent transcription of the gene in target cells. See generally, Helene. (1991) *Anticancer Drug Des.* 6: 569-84; Helene. *et al.* (1992) *Ann. N.Y. Acad. Sci.* 660:27-36; and Maher (1992) *Bioassays* 14: 807-15.

In various embodiments, the nucleic acids of the invention can be modified at the base moiety, sugar moiety or phosphate backbone to improve, *e.g.*, the stability, hybridization, or solubility of the molecule. For example, the deoxyribose phosphate backbone of the nucleic acids can be modified to generate peptide nucleic acids (see Hyrup *et al.* (1996) *Bioorg Med Chem* 4: 5-23). As used herein, the terms "peptide nucleic acids" or "PNAs" refer to nucleic acid mimics, *e.g.*, DNA mimics, in which the deoxyribose phosphate backbone is replaced by a pseudopeptide backbone and only the four natural nucleobases are retained. The neutral backbone of PNAs has been shown to allow for specific hybridization to DNA and RNA under conditions of low ionic strength. The synthesis of PNA oligomers can be performed using standard solid phase peptide synthesis protocols as described in Hyrup *et al.* (1996) above; Perry-O'Keefe *et al.* (1996) *PNAS* 93: 14670-675.

PNAs of the invention can be used in therapeutic and diagnostic applications. For example, PNAs can be used as antisense or antigene agents for sequence-specific modulation of gene expression by, *e.g.*, inducing transcription or translation arrest or inhibiting replication. PNAs of the invention can also be used, *e.g.*, in the analysis of single base pair mutations in a gene by, *e.g.*, PNA directed PCR clamping; as artificial restriction enzymes when used in combination with other enzymes, *e.g.*, S1 nucleases (Hyrup B. (1996) above); or as probes or primers for DNA sequence and hybridization (Hyrup *et al.* (1996), above; Perry-O'Keefe (1996), above).

In another embodiment, PNAs of the invention can be modified, *e.g.*, to enhance their stability or cellular uptake, by attaching lipophilic or other helper groups to PNA, by the formation of PNA-DNA chimeras, or by the use of liposomes or other techniques of drug delivery known in the art. For example, PNA-DNA chimeras can be generated that may

combine the advantageous properties of PNA and DNA. Such chimeras allow DNA recognition enzymes, *e.g.*, RNase H and DNA polymerases, to interact with the DNA portion while the PNA portion would provide high binding affinity and specificity. PNA-DNA chimeras can be linked using linkers of appropriate lengths selected in terms of base stacking, number of bonds between the nucleobases, and orientation (Hyrup (1996) above). The synthesis of PNA-DNA chimeras can be performed as described in Hyrup (1996) above and Finn *et al.* (1996) *Nucl Acids Res* 24: 3357-63. For example, a DNA chain can be synthesized on a solid support using standard phosphoramidite coupling chemistry, and modified nucleoside analogs, *e.g.*, 5'-(4-methoxytrityl)amino-5'-deoxy-thymidine phosphoramidite, can be used between the PNA and the 5' end of DNA (Mag *et al.* (1989) *Nucl Acid Res* 17: 5973-88). PNA monomers are then coupled in a stepwise manner to produce a chimeric molecule with a 5' PNA segment and a 3' DNA segment (Finn *et al.* (1996) above). Alternatively, chimeric molecules can be synthesized with a 5' DNA segment and a 3' PNA segment. See, Petersen *et al.* (1975) *Bioorg Med Chem Lett* 5: 1119-11124.

In other embodiments, the oligonucleotide may include other appended groups such as peptides (*e.g.*, for targeting host cell receptors *in vivo*), or agents facilitating transport across the cell membrane (see, *e.g.*, Letsinger *et al.*, 1989, *Proc. Natl. Acad. Sci. U.S.A.* 86:6553-6556; Lemaitre *et al.*, 1987, Proc. *Natl. Acad. Sci.* 84:648-652; PCT Publication No. W088/09810) or the blood-brain barrier (see, *e.g.*, PCT Publication No. W089/10134). In addition, oligonucleotides can be modified with hybridization triggered cleavage agents (See, *e.g.*, Krol *et al.*, 1988, *BioTechniques* 6:958-976) or intercalating agents. (See, *e.g.*, Zon, 1988, *Pharm. Res.* 5: 539-549). To this end, the oligonucleotide may be conjugated to another molecule, *e.g.*, a peptide, a hybridization triggered cross-linking agent, a transport agent, a hybridization-triggered cleavage agent, etc.

## 4.5 HOSTS

The present invention further provides host cells genetically engineered to contain the polynucleotides of the invention. For example, such host cells may contain nucleic acids of the invention introduced into the host cell using known transformation, transfection or infection methods. The present invention still further provides host cells genetically engineered to express the polynucleotides of the invention, wherein such polynucleotides are in operative association with a regulatory sequence heterologous to the host cell which drives expression of the polynucleotides in the cell.

Knowledge of nucleic acid sequences allows for modification of cells to permit, or increase, expression of endogenous polypeptide. Cells can be modified (*e.g.*, by homologous

recombination) to provide increased polypeptide expression by replacing, in whole or in part, the naturally occurring promoter with all or part of a heterologous promoter so that the cells express the polypeptide at higher levels. The heterologous promoter is inserted in such a manner that it is operatively linked to the encoding sequences. See, for example, PCT International Publication No. WO94/12650, PCT International Publication No. WO92/20808, and PCT International Publication No. WO91/09955. It is also contemplated that, in addition to heterologous promoter DNA, amplifiable marker DNA (*e.g.*, ada, dhfr, and the multifunctional CAD gene which encodes carbamyl phosphate synthase, aspartate transcarbamylase, and dihydroorotase) and/or intron DNA may be inserted along with the heterologous promoter DNA. If linked to the coding sequence, amplification of the marker DNA by standard selection methods results in co-amplification of the desired protein coding sequences in the cells.

The host cell can be a higher eukaryotic host cell, such as a mammalian cell, a lower eukaryotic host cell, such as a yeast cell, or the host cell can be a prokaryotic cell, such as a bacterial cell. Introduction of the recombinant construct into the host cell can be effected by calcium phosphate transfection, DEAE, dextran mediated transfection, or electroporation (Davis, L. et al., *Basic Methods in Molecular Biology* (1986)). The host cells containing one of the polynucleotides of the invention, can be used in conventional manners to produce the gene product encoded by the isolated fragment (in the case of an ORF) or can be used to produce a heterologous protein under the control of the EMF.

Any host/vector system can be used to express one or more of the ORFs of the present invention. These include, but are not limited to, eukaryotic hosts such as HeLa cells, Cv-1 cell, COS cells, 293 cells, and Sf9 cells, as well as prokaryotic host such as *E. coli* and *B. subtilis.* The most preferred cells are those which do not normally express the particular polypeptide or protein or which expresses the polypeptide or protein at low natural level. Mature proteins can be expressed in mammalian cells, yeast, bacteria, or other cells under the control of appropriate promoters. Cell-free translation systems can also be employed to produce such proteins using RNAs derived from the DNA constructs of the present invention. Appropriate cloning and expression vectors for use with prokaryotic and eukaryotic hosts are described by Sambrook, et al., in Molecular Cloning: A Laboratory Manual, Second Edition, Cold Spring Harbor, New York (1989), the disclosure of which is hereby incorporated by reference.

Various mammalian cell culture systems can also be employed to express recombinant protein. Examples of mammalian expression systems include the COS-7 lines of monkey kidney fibroblasts, described by Gluzman, Cell 23:175 (1981). Other cell lines capable of expressing a compatible vector are, for example, the C127, monkey COS cells, Chinese Hamster Ovary (CHO) cells, human kidney 293 cells, human epidermal A431 cells, human Colo205 cells, 3T3

cells, CV-1 cells, other transformed primate cell lines, normal diploid cells, cell strains derived from *in vitro* culture of primary tissue, primary explants, HeLa cells, mouse L cells, BHK, HL-60, U937, HaK or Jurkat cells. Mammalian expression vectors will comprise an origin of replication, a suitable promoter and also any necessary ribosome binding sites, polyadenylation site, splice donor and acceptor sites, transcriptional termination sequences, and 5′ flanking nontranscribed sequences. DNA sequences derived from the SV40 viral genome, for example, SV40 origin, early promoter, enhancer, splice, and polyadenylation sites may be used to provide the required nontranscribed genetic elements. Recombinant polypeptides and proteins produced in bacterial culture are usually isolated by initial extraction from cell pellets, followed by one or more salting-out, aqueous ion exchange or size exclusion chromatography steps. Protein refolding steps can be used, as necessary, in completing configuration of the mature protein. Finally, high performance liquid chromatography (HPLC) can be employed for final purification steps. Microbial cells employed in expression of proteins can be disrupted by any convenient method, including freeze-thaw cycling, sonication, mechanical disruption, or use of cell lysing agents.

Alternatively, it may be possible to produce the protein in lower eukaryotes such as yeast or insects or in prokaryotes such as bacteria. Potentially suitable yeast strains include *Saccharomyces cerevisiae*, *Schizosaccharomyces pombe*, *Kluyveromyces* strains, *Candida*, or any yeast strain capable of expressing heterologous proteins. Potentially suitable bacterial strains include *Escherichia coli, Bacillus subtilis, Salmonella typhimurium*, or any bacterial strain capable of expressing heterologous proteins. If the protein is made in yeast or bacteria, it may be necessary to modify the protein produced therein, for example by phosphorylation or glycosylation of the appropriate sites, in order to obtain the functional protein. Such covalent attachments may be accomplished using known chemical or enzymatic methods.

In another embodiment of the present invention, cells and tissues may be engineered to express an endogenous gene comprising the polynucleotides of the invention under the control of inducible regulatory elements, in which case the regulatory sequences of the endogenous gene may be replaced by homologous recombination. As described herein, gene targeting can be used to replace a gene's existing regulatory region with a regulatory sequence isolated from a different gene or a novel regulatory sequence synthesized by genetic engineering methods. Such regulatory sequences may be comprised of promoters, enhancers, scaffold-attachment regions, negative regulatory elements, transcriptional initiation sites, regulatory protein binding sites or combinations of said sequences. Alternatively, sequences which affect the structure or stability of the RNA or protein produced may be replaced, removed, added, or otherwise modified by targeting. These sequence include polyadenylation signals, mRNA stability elements, splice

sites, leader sequences for enhancing or modifying transport or secretion properties of the protein, or other sequences which alter or improve the function or stability of protein or RNA molecules.

The targeting event may be a simple insertion of the regulatory sequence, placing the gene under the control of the new regulatory sequence, *e.g.*, inserting a new promoter or enhancer or both upstream of a gene. Alternatively, the targeting event may be a simple deletion of a regulatory element, such as the deletion of a tissue-specific negative regulatory element. Alternatively, the targeting event may replace an existing element; for example, a tissue-specific enhancer can be replaced by an enhancer that has broader or different cell-type specificity than the naturally occurring elements. Here, the naturally occurring sequences are deleted and new sequences are added. In all cases, the identification of the targeting event may be facilitated by the use of one or more selectable marker genes that are contiguous with the targeting DNA, allowing for the selection of cells in which the exogenous DNA has integrated into the host cell genome. The identification of the targeting event may also be facilitated by the use of one or more marker genes exhibiting the property of negative selection, such that the negatively selectable marker is linked to the exogenous DNA, but configured such that the negatively selectable marker flanks the targeting sequence, and such that a correct homologous recombination event with sequences in the host cell genome does not result in the stable integration of the negatively selectable marker. Markers useful for this purpose include the Herpes Simplex Virus thymidine kinase (TK) gene or the bacterial xanthine-guanine phosphoribosyl-transferase (gpt) gene.

The gene targeting or gene activation techniques which can be used in accordance with this aspect of the invention are more particularly described in U.S. Patent No. 5,272,071 to Chappel; U.S. Patent No. 5,578,461 to Sherwin et al.; International Application No. PCT/US92/09627 (WO93/09222) by Selden et al.; and International Application No. PCT/US90/06436 (WO91/06667) by Skoultchi et al., each of which is incorporated by reference herein in its entirety.

### 4.6 POLYPEPTIDES OF THE INVENTION

The isolated polypeptides of the invention include, but are not limited to, a polypeptide comprising: the amino acid sequences set forth as any one of SEQ ID NO:1010-2018 or an amino acid sequence encoded by any one of the nucleotide sequences SEQ ID NO:1-1009 or the corresponding full length or mature protein. Polypeptides of the invention also include polypeptides preferably with biological or immunological activity that are encoded by: (a) a polynucleotide having any one of the nucleotide sequences set forth in SEQ ID NO:1-1009 or (b)

polynucleotides encoding any one of the amino acid sequences set forth as SEQ ID NO:1010-2018 or (c) polynucleotides that hybridize to the complement of the polynucleotides of either (a) or (b) under stringent hybridization conditions. The invention also provides biologically active or immunologically active variants of any of the amino acid sequences set forth as SEQ ID NO:1010-2018 or the corresponding full length or mature protein; and "substantial equivalents" thereof (*e.g.*, with at least about 65%, at least about 70%, at least about 75%, at least about 80%, at least about 85%, 86%, 87%, 88%, 89%, at least about 90%, 91%, 92%, 93%, 94%, typically at least about 95%, 96%, 97%, more typically at least about 98%, or most typically at least about 99% amino acid identity) that retain biological activity. Polypeptides encoded by allelic variants may have a similar, increased, or decreased activity compared to polypeptides comprising SEQ ID NO:1010-2018.

Fragments of the proteins of the present invention which are capable of exhibiting biological activity are also encompassed by the present invention. Fragments of the protein may be in linear form or they may be cyclized using known methods, for example, as described in H. U. Saragovi, et al., Bio/Technology 10, 773-778 (1992) and in R. S. McDowell, et al., J. Amer. Chem. Soc. 114, 9245-9253 (1992), both of which are incorporated herein by reference. Such fragments may be fused to carrier molecules such as immunoglobulins for many purposes, including increasing the valency of protein binding sites.

The present invention also provides both full-length and mature forms (for example, without a signal sequence or precursor sequence) of the disclosed proteins. The protein coding sequence is identified in the sequence listing by translation of the disclosed nucleotide sequences. The mature form of such protein may be obtained by expression of a full-length polynucleotide in a suitable mammalian cell or other host cell. The sequence of the mature form of the protein is also determinable from the amino acid sequence of the full-length form. Where proteins of the present invention are membrane bound, soluble forms of the proteins are also provided. In such forms, part or all of the regions causing the proteins to be membrane bound are deleted so that the proteins are fully secreted from the cell in which they are expressed.

Protein compositions of the present invention may further comprise an acceptable carrier, such as a hydrophilic, *e.g.*, pharmaceutically acceptable, carrier.

The present invention further provides isolated polypeptides encoded by the nucleic acid fragments of the present invention or by degenerate variants of the nucleic acid fragments of the present invention. By "degenerate variant" is intended nucleotide fragments which differ from a nucleic acid fragment of the present invention (*e.g.*, an ORF) by nucleotide sequence but, due to the degeneracy of the genetic code, encode an identical polypeptide sequence. Preferred nucleic acid fragments of the present invention are the ORFs that encode proteins.

A variety of methodologies known in the art can be utilized to obtain any one of the isolated polypeptides or proteins of the present invention. At the simplest level, the amino acid sequence can be synthesized using commercially available peptide synthesizers. The synthetically-constructed protein sequences, by virtue of sharing primary, secondary or tertiary structural and/or conformational characteristics with proteins may possess biological properties in common therewith, including protein activity. This technique is particularly useful in producing small peptides and fragments of larger polypeptides. Fragments are useful, for example, in generating antibodies against the native polypeptide. Thus, they may be employed as biologically active or immunological substitutes for natural, purified proteins in screening of therapeutic compounds and in immunological processes for the development of antibodies.

The polypeptides and proteins of the present invention can alternatively be purified from cells which have been altered to express the desired polypeptide or protein. As used herein, a cell is said to be altered to express a desired polypeptide or protein when the cell, through genetic manipulation, is made to produce a polypeptide or protein which it normally does not produce or which the cell normally produces at a lower level. One skilled in the art can readily adapt procedures for introducing and expressing either recombinant or synthetic sequences into eukaryotic or prokaryotic cells in order to generate a cell which produces one of the polypeptides or proteins of the present invention.

The invention also relates to methods for producing a polypeptide comprising growing a culture of host cells of the invention in a suitable culture medium, and purifying the protein from the cells or the culture in which the cells are grown. For example, the methods of the invention include a process for producing a polypeptide in which a host cell containing a suitable expression vector that includes a polynucleotide of the invention is cultured under conditions that allow expression of the encoded polypeptide. The polypeptide can be recovered from the culture, conveniently from the culture medium, or from a lysate prepared from the host cells and further purified. Preferred embodiments include those in which the protein produced by such process is a full length or mature form of the protein.

In an alternative method, the polypeptide or protein is purified from bacterial cells which naturally produce the polypeptide or protein. One skilled in the art can readily follow known methods for isolating polypeptides and proteins in order to obtain one of the isolated polypeptides or proteins of the present invention. These include, but are not limited to, immunochromatography, HPLC, size-exclusion chromatography, ion-exchange chromatography, and immuno-affinity chromatography. See, *e.g.*, Scopes, *Protein Purification: Principles and Practice*, Springer-Verlag (1994); Sambrook, et al., in Molecular Cloning: *A Laboratory Manual*; Ausubel et al., *Current Protocols in Molecular Biology*. Polypeptide fragments that

retain biological/immunological activity include fragments comprising greater than about 100 amino acids, or greater than about 200 amino acids, and fragments that encode specific protein domains.

The purified polypeptides can be used in *in vitro* binding assays which are well known in the art to identify molecules which bind to the polypeptides. These molecules include but are not limited to, for *e.g.*, small molecules, molecules from combinatorial libraries, antibodies or other proteins. The molecules identified in the binding assay are then tested for antagonist or agonist activity in *in vivo* tissue culture or animal models that are well known in the art. In brief, the molecules are titrated into a plurality of cell cultures or animals and then tested for either cell/animal death or prolonged survival of the animal/cells.

In addition, the peptides of the invention or molecules capable of binding to the peptides may be complexed with toxins, *e.g.*, ricin or cholera, or with other compounds that are toxic to cells. The toxin-binding molecule complex is then targeted to a tumor or other cell by the specificity of the binding molecule for SEQ ID NO:1010-2018.

The protein of the invention may also be expressed as a product of transgenic animals, *e.g.*, as a component of the milk of transgenic cows, goats, pigs, or sheep which are characterized by somatic or germ cells containing a nucleotide sequence encoding the protein.

The proteins provided herein also include proteins characterized by amino acid sequences similar to those of purified proteins but into which modification are naturally provided or deliberately engineered. For example, modifications, in the peptide or DNA sequence, can be made by those skilled in the art using known techniques. Modifications of interest in the protein sequences may include the alteration, substitution, replacement, insertion or deletion of a selected amino acid residue in the coding sequence. For example, one or more of the cysteine residues may be deleted or replaced with another amino acid to alter the conformation of the molecule. Techniques for such alteration, substitution, replacement, insertion or deletion are well known to those skilled in the art (see, *e.g.*, U.S. Pat. No. 4,518,584). Preferably, such alteration, substitution, replacement, insertion or deletion retains the desired activity of the protein. Regions of the protein that are important for the protein function can be determined by various methods known in the art including the alanine-scanning method which involved systematic substitution of single or strings of amino acids with alanine, followed by testing the resulting alanine-containing variant for biological activity. This type of analysis determines the importance of the substituted amino acid(s) in biological activity. Regions of the protein that are important for protein function may be determined by the eMATRIX program.

Other fragments and derivatives of the sequences of proteins which would be expected to retain protein activity in whole or in part and are useful for screening or other immunological

methodologies may also be easily made by those skilled in the art given the disclosures herein. Such modifications are encompassed by the present invention.

The protein may also be produced by operably linking the isolated polynucleotide of the invention to suitable control sequences in one or more insect expression vectors, and employing an insect expression system. Materials and methods for baculovirus/insect cell expression systems are commercially available in kit form from, *e.g.*, Invitrogen, San Diego, Calif., U.S.A. (the MaxBat™ kit), and such methods are well known in the art, as described in Summers and Smith, Texas Agricultural Experiment Station Bulletin No. 1555 (1987), incorporated herein by reference. As used herein, an insect cell capable of expressing a polynucleotide of the present invention is "transformed."

The protein of the invention may be prepared by culturing transformed host cells under culture conditions suitable to express the recombinant protein. The resulting expressed protein may then be purified from such culture (*i.e.*, from culture medium or cell extracts) using known purification processes, such as gel filtration and ion exchange chromatography. The purification of the protein may also include an affinity column containing agents which will bind to the protein; one or more column steps over such affinity resins as concanavalin A-agarose, heparin-toyopearl™ or Cibacrom blue 3GA Sepharose™; one or more steps involving hydrophobic interaction chromatography using such resins as phenyl ether, butyl ether, or propyl ether; or immunoaffinity chromatography.

Alternatively, the protein of the invention may also be expressed in a form which will facilitate purification. For example, it may be expressed as a fusion protein, such as those of maltose binding protein (MBP), glutathione-S-transferase (GST) or thioredoxin (TRX), or as a His tag. Kits for expression and purification of such fusion proteins are commercially available from New England BioLab (Beverly, Mass.), Pharmacia (Piscataway, N.J.) and Invitrogen, respectively. The protein can also be tagged with an epitope and subsequently purified by using a specific antibody directed to such epitope. One such epitope ("FLAG®") is commercially available from Kodak (New Haven, Conn.).

Finally, one or more reverse-phase high performance liquid chromatography (RP- HPLC) steps employing hydrophobic RP-HPLC media, *e.g.*, silica gel having pendant methyl or other aliphatic groups, can be employed to further purify the protein. Some or all of the foregoing purification steps, in various combinations, can also be employed to provide a substantially homogeneous isolated recombinant protein. The protein thus purified is substantially free of other mammalian proteins and is defined in accordance with the present invention as an "isolated protein."

The polypeptides of the invention include analogs (variants). This embraces fragments, as well as peptides in which one or more amino acids has been deleted, inserted, or substituted. Also, analogs of the polypeptides of the invention embrace fusions of the polypeptides or modifications of the polypeptides of the invention, wherein the polypeptide or analog is fused to another moiety or moieties, *e.g.*, targeting moiety or another therapeutic agent. Such analogs may exhibit improved properties such as activity and/or stability. Examples of moieties which may be fused to the polypeptide or an analog include, for example, targeting moieties which provide for the delivery of polypeptide to pancreatic cells, *e.g.*, antibodies to pancreatic cells, antibodies to immune cells such as T-cells, monocytes, dendritic cells, granulocytes, etc., as well as receptor and ligands expressed on pancreatic or immune cells. Other moieties which may be fused to the polypeptide include therapeutic agents which are used for treatment, for example, immunosuppressive drugs such as cyclosporin, SK506, azathioprine, CD3 antibodies and steroids. Also, polypeptides may be fused to immune modulators, and other cytokines such as alpha or beta interferon.

### 4.6.1 DETERMINING POLYPEPTIDE AND POLYNUCLEOTIDE IDENTITY AND SIMILARITY

Preferred identity and/or similarity are designed to give the largest match between the sequences tested. Methods to determine identity and similarity are codified in computer programs including, but are not limited to, the GCG program package, including GAP (Devereux, J., et al., Nucleic Acids Research 12(1):387 (1984); Genetics Computer Group, University of Wisconsin, Madison, WI), BLASTP, BLASTN, BLASTX, FASTA (Altschul, S.F. et al., J. Molec. Biol. 215:403-410 (1990), PSI-BLAST (Altschul S.F. et al., Nucleic Acids Res. vol. 25, pp. 3389-3402, herein incorporated by reference), eMatrix software (Wu et al., J. Comp. Biol., Vol. 6, pp. 219-235 (1999), herein incorporated by reference), eMotif software (Nevill-Manning et al, ISMB-97, Vol. 4, pp. 202-209, herein incorporated by reference), pFam software (Sonnhammer et al., Nucleic Acids Res., Vol. 26(1), pp. 320-322 (1998), herein incorporated by reference) and the Kyte-Doolittle hydrophobocity prediction algorithm (J. Mol Biol, 157, pp. 105-31 (1982), incorporated herein by reference). The BLAST programs are publicly available from the National Center for Biotechnology Information (NCBI) and other sources (BLAST Manual, Altschul, S., et al. NCB NLM NIH Bethesda, MD 20894; Altschul, S., et al., J. Mol. Biol. 215:403-410 (1990).

### 4.7 CHIMERIC AND FUSION PROTEINS

The invention also provides chimeric or fusion proteins. As used herein, a "chimeric protein" or "fusion protein" comprises a polypeptide of the invention operatively linked to

another polypeptide. Within a fusion protein the polypeptide according to the invention can correspond to all or a portion of a protein according to the invention. In one embodiment, a fusion protein comprises at least one biologically active portion of a protein according to the invention. In another embodiment, a fusion protein comprises at least two biologically active portions of a protein according to the invention. Within the fusion protein, the term "operatively linked" is intended to indicate that the polypeptide according to the invention and the other polypeptide are fused in-frame to each other. The polypeptide can be fused to the N-terminus or C-terminus.

For example, in one embodiment a fusion protein comprises a polypeptide according to the invention operably linked to the extracellular domain of a second protein.

In another embodiment, the fusion protein is a GST-fusion protein in which the polypeptide sequences of the invention are fused to the C-terminus of the GST (*i.e.*, glutathione S-transferase) sequences.

In another embodiment, the fusion protein is an immunoglobulin fusion protein in which the polypeptide sequences according to the invention comprises one or more domains are fused to sequences derived from a member of the immunoglobulin protein family. The immunoglobulin fusion proteins of the invention can be incorporated into pharmaceutical compositions and administered to a subject to inhibit an interaction between a ligand and a protein of the invention on the surface of a cell, to thereby suppress signal transduction *in vivo*. The immunoglobulin fusion proteins can be used to affect the bioavailability of a cognate ligand. Inhibition of the ligand/protein interaction may be useful therapeutically for both the treatment of proliferative and differentiative disorders, *e,g.,* cancer as well as modulating (*e.g.*, promoting or inhibiting) cell survival. Moreover, the immunoglobulin fusion proteins of the invention can be used as immunogens to produce antibodies in a subject, to purify ligands, and in screening assays to identify molecules that inhibit the interaction of a polypeptide of the invention with a ligand.

A chimeric or fusion protein of the invention can be produced by standard recombinant DNA techniques. For example, DNA fragments coding for the different polypeptide sequences are ligated together in-frame in accordance with conventional techniques, *e.g.*, by employing blunt-ended or stagger-ended termini for ligation, restriction enzyme digestion to provide for appropriate termini, filling-in of cohesive ends as appropriate, alkaline phosphatase treatment to avoid undesirable joining, and enzymatic ligation. In another embodiment, the fusion gene can be synthesized by conventional techniques including automated DNA synthesizers. Alternatively, PCR amplification of gene fragments can be carried out using anchor primers that give rise to complementary overhangs between two consecutive gene fragments that can subsequently be annealed and reamplified to generate a chimeric gene sequence (see, for

example, Ausubel et al. (eds.) CURRENT PROTOCOLS IN MOLECULAR BIOLOGY, John Wiley & Sons, 1992). Moreover, many expression vectors are commercially available that already encode a fusion moiety (*e.g.*, a GST polypeptide). A nucleic acid encoding a polypeptide of the invention can be cloned into such an expression vector such that the fusion moiety is linked in-frame to the protein of the invention.

## 4.8 GENE THERAPY

Mutations in the polynucleotides of the invention gene may result in loss of normal function of the encoded protein. The invention thus provides gene therapy to restore normal activity of the polypeptides of the invention; or to treat disease states involving polypeptides of the invention. Delivery of a functional gene encoding polypeptides of the invention to appropriate cells is effected *ex vivo*, *in situ*, or *in vivo* by use of vectors, and more particularly viral vectors (*e.g.*, adenovirus, adeno-associated virus, or a retrovirus), or *ex vivo* by use of physical DNA transfer methods (*e.g.*, liposomes or chemical treatments). See, for example, Anderson, Nature, supplement to vol. 392, no. 6679, pp.25-20 (1998). For additional reviews of gene therapy technology see Friedmann, Science, 244: 1275-1281 (1989); Verma, Scientific American: 68-84 (1990); and Miller, Nature, 357: 455-460 (1992). Introduction of any one of the nucleotides of the present invention or a gene encoding the polypeptides of the present invention can also be accomplished with extrachromosomal substrates (transient expression) or artificial chromosomes (stable expression). Cells may also be cultured *ex vivo* in the presence of proteins of the present invention in order to proliferate or to produce a desired effect on or activity in such cells. Treated cells can then be introduced *in vivo* for therapeutic purposes. Alternatively, it is contemplated that in other human disease states, preventing the expression of or inhibiting the activity of polypeptides of the invention will be useful in treating the disease states. It is contemplated that antisense therapy or gene therapy could be applied to negatively regulate the expression of polypeptides of the invention.

Other methods inhibiting expression of a protein include the introduction of antisense molecules to the nucleic acids of the present invention~~, their complements~~, or their ~~translated~~ RNA sequences, by methods known in the art. Further, the polypeptides of the present invention can be inhibited by using targeted deletion methods, or the insertion of a negative regulatory element such as a silencer, which is tissue specific.

The present invention still further provides cells genetically engineered *in vivo* to express the polynucleotides of the invention, wherein such polynucleotides are in operative association with a regulatory sequence heterologous to the host cell which drives expression of the polynucleotides in

the cell. These methods can be used to increase or decrease the expression of the polynucleotides of the present invention.

Knowledge of DNA sequences provided by the invention allows for modification of cells to permit, increase, or decrease, expression of endogenous polypeptide. Cells can be modified (*e.g.*, by homologous recombination) to provide increased polypeptide expression by replacing, in whole or in part, the naturally occurring promoter with all or part of a heterologous promoter so that the cells express the protein at higher levels. The heterologous promoter is inserted in such a manner that it is operatively linked to the desired protein encoding sequences. See, for example, PCT International Publication No. WO 94/12650, PCT International Publication No. WO 92/20808, and PCT International Publication No. WO 91/09955. It is also contemplated that, in addition to heterologous promoter DNA, amplifiable marker DNA (*e.g.*, ada, dhfr, and the multifunctional CAD gene which encodes carbamyl phosphate synthase, aspartate transcarbamylase, and dihydroorotase) and/or intron DNA may be inserted along with the heterologous promoter DNA. If linked to the desired protein coding sequence, amplification of the marker DNA by standard selection methods results in co-amplification of the desired protein coding sequences in the cells.

In another embodiment of the present invention, cells and tissues may be engineered to express an endogenous gene comprising the polynucleotides of the invention under the control of inducible regulatory elements, in which case the regulatory sequences of the endogenous gene may be replaced by homologous recombination. As described herein, gene targeting can be used to replace a gene's existing regulatory region with a regulatory sequence isolated from a different gene or a novel regulatory sequence synthesized by genetic engineering methods. Such regulatory sequences may be comprised of promoters, enhancers, scaffold-attachment regions, negative regulatory elements, transcriptional initiation sites, regulatory protein binding sites or combinations of said sequences. Alternatively, sequences which affect the structure or stability of the RNA or protein produced may be replaced, removed, added, or otherwise modified by targeting. These sequences include polyadenylation signals, mRNA stability elements, splice sites, leader sequences for enhancing or modifying transport or secretion properties of the protein, or other sequences which alter or improve the function or stability of protein or RNA molecules.

The targeting event may be a simple insertion of the regulatory sequence, placing the gene under the control of the new regulatory sequence, *e.g.,* inserting a new promoter or enhancer or both upstream of a gene. Alternatively, the targeting event may be a simple deletion of a regulatory element, such as the deletion of a tissue-specific negative regulatory element. Alternatively, the targeting event may replace an existing element; for example, a tissue-specific enhancer can be replaced by an enhancer that has broader or different cell-type specificity than the naturally occurring elements. Here, the naturally occurring sequences are deleted and new sequences are

added. In all cases, the identification of the targeting event may be facilitated by the use of one or more selectable marker genes that are contiguous with the targeting DNA, allowing for the selection of cells in which the exogenous DNA has integrated into the cell genome. The identification of the targeting event may also be facilitated by the use of one or more marker genes exhibiting the property of negative selection, such that the negatively selectable marker is linked to the exogenous DNA, but configured such that the negatively selectable marker flanks the targeting sequence, and such that a correct homologous recombination event with sequences in the host cell genome does not result in the stable integration of the negatively selectable marker. Markers useful for this purpose include the Herpes Simplex Virus thymidine kinase (TK) gene or the bacterial xanthine-guanine phosphoribosyl-transferase (gpt) gene.

The gene targeting or gene activation techniques which can be used in accordance with this aspect of the invention are more particularly described in U.S. Patent No. 5,272,071 to Chappel; U.S. Patent No. 5,578,461 to Sherwin et al.; International Application No. PCT/US92/09627 (WO93/09222) by Selden et al.; and International Application No. PCT/US90/06436 (WO91/06667) by Skoultchi et al., each of which is incorporated by reference herein in its entirety.

## 4.9 TRANSGENIC ANIMALS

In preferred methods to determine biological functions of the polypeptides of the invention in vivo, one or more genes provided by the invention are either over expressed or inactivated in the germ line of animals using homologous recombination [Capecchi, Science 244:1288-1292 (1989)]. Animals in which the gene is over expressed, under the regulatory control of exogenous or endogenous promoter elements, are known as transgenic animals. Animals in which an endogenous gene has been inactivated by homologous recombination are referred to as "knockout" animals. Knockout animals, preferably non-human mammals, can be prepared as described in U.S. Patent No. 5,557,032, incorporated herein by reference. Transgenic animals are useful to determine the roles polypeptides of the invention play in biological processes, and preferably in disease states. Transgenic animals are useful as model systems to identify compounds that modulate lipid metabolism. Transgenic animals, preferably non-human mammals, are produced using methods as described in U.S. Patent No 5,489,743 and PCT Publication No. WO94/28122, incorporated herein by reference.

Transgenic animals can be prepared wherein all or part of a promoter of the polynucleotides of the invention is either activated or inactivated to alter the level of expression of the polypeptides of the invention. Inactivation can be carried out using homologous recombination methods described above. Activation can be achieved by supplementing or even replacing the homologous promoter to provide for increased protein expression. The homologous

promoter can be supplemented by insertion of one or more heterologous enhancer elements known to confer promoter activation in a particular tissue.

The polynucleotides of the present invention also make possible the development, through, *e.g.*, homologous recombination or knock out strategies, of animals that fail to express polypeptides of the invention or that express a variant polypeptide. Such animals are useful as models for studying the *in vivo* activities of polypeptide as well as for studying modulators of the polypeptides of the invention.

In preferred methods to determine biological functions of the polypeptides of the invention *in vivo,* one or more genes provided by the invention are either over expressed or inactivated in the germ line of animals using homologous recombination [Capecchi, Science 244:1288-1292 (1989)]. Animals in which the gene is over expressed, under the regulatory control of exogenous or endogenous promoter elements, are known as transgenic animals. Animals in which an endogenous gene has been inactivated by homologous recombination are referred to as "knockout" animals. Knockout animals, preferably non-human mammals, can be prepared as described in U.S. Patent No. 5,557,032, incorporated herein by reference. Transgenic animals are useful to determine the roles polypeptides of the invention play in biological processes, and preferably in disease states. Transgenic animals are useful as model systems to identify compounds that modulate lipid metabolism. Transgenic animals, preferably non-human mammals, are produced using methods as described in U.S. Patent No 5,489,743 and PCT Publication No. WO94/28122, incorporated herein by reference.

Transgenic animals can be prepared wherein all or part of the polynucleotides of the invention promoter is either activated or inactivated to alter the level of expression of the polypeptides of the invention. Inactivation can be carried out using homologous recombination methods described above. Activation can be achieved by supplementing or even replacing the homologous promoter to provide for increased protein expression. The homologous promoter can be supplemented by insertion of one or more heterologous enhancer elements known to confer promoter activation in a particular tissue.

## 4.10 USES AND BIOLOGICAL ACTIVITY

The polynucleotides and proteins of the present invention are expected to exhibit one or more of the uses or biological activities (including those associated with assays cited herein) identified herein. Uses or activities described for proteins of the present invention may be provided by administration or use of such proteins or of polynucleotides encoding such proteins (such as, for example, in gene therapies or vectors suitable for introduction of DNA). The mechanism underlying the particular condition or pathology will dictate whether the

polypeptides of the invention, the polynucleotides of the invention or modulators (activators or inhibitors) thereof would be beneficial to the subject in need of treatment. Thus, "therapeutic compositions of the invention" include compositions comprising isolated polynucleotides (including recombinant DNA molecules, cloned genes and degenerate variants thereof) or polypeptides of the invention (including full length protein, mature protein and truncations or domains thereof), or compounds and other substances that modulate the overall activity of the target gene products, either at the level of target gene/protein expression or target protein activity. Such modulators include polypeptides, analogs, (variants), including fragments and fusion proteins, antibodies and other binding proteins; chemical compounds that directly or indirectly activate or inhibit the polypeptides of the invention (identified, *e.g.*, via drug screening assays as described herein); antisense polynucleotides and polynucleotides suitable for triple helix formation; and in particular antibodies or other binding partners that specifically recognize one or more epitopes of the polypeptides of the invention.

The polypeptides of the present invention may likewise be involved in cellular activation or in one of the other physiological pathways described herein.

### 4.10.1 RESEARCH USES AND UTILITIES

The polynucleotides provided by the present invention can be used by the research community for various purposes. The polynucleotides can be used to express recombinant protein for analysis, characterization or therapeutic use; as markers for tissues in which the corresponding protein is preferentially expressed (either constitutively or at a particular stage of tissue differentiation or development or in disease states); as molecular weight markers on gels; as chromosome markers or tags (when labeled) to identify chromosomes or to map related gene positions; to compare with endogenous DNA sequences in patients to identify potential genetic disorders; as probes to hybridize and thus discover novel, related DNA sequences; as a source of information to derive PCR primers for genetic fingerprinting; as a probe to "subtract-out" known sequences in the process of discovering other novel polynucleotides; for selecting and making oligomers for attachment to a "gene chip" or other support, including for examination of expression patterns; to raise anti-protein antibodies using DNA immunization techniques; and as an antigen to raise anti-DNA antibodies or elicit another immune response. Where the polynucleotide encodes a protein which binds or potentially binds to another protein (such as, for example, in a receptor-ligand interaction), the polynucleotide can also be used in interaction trap assays (such as, for example, that described in Gyuris et al., Cell 75:791-803 (1993)) to identify polynucleotides encoding the other protein with which binding occurs or to identify inhibitors of the binding interaction.

The polypeptides provided by the present invention can similarly be used in assays to determine biological activity, including in a panel of multiple proteins for high-throughput screening; to raise antibodies or to elicit another immune response; as a reagent (including the labeled reagent) in assays designed to quantitatively determine levels of the protein (or its receptor) in biological fluids; as markers for tissues in which the corresponding polypeptide is preferentially expressed (either constitutively or at a particular stage of tissue differentiation or development or in a disease state); and, of course, to isolate correlative receptors or ligands. Proteins involved in these binding interactions can also be used to screen for peptide or small molecule inhibitors or agonists of the binding interaction.

Any or all of these research utilities are capable of being developed into reagent grade or kit format for commercialization as research products.

Methods for performing the uses listed above are well known to those skilled in the art. References disclosing such methods include without limitation "Molecular Cloning: A Laboratory Manual", 2d ed., Cold Spring Harbor Laboratory Press, Sambrook, J., E. F. Fritsch and T. Maniatis eds., 1989, and "Methods in Enzymology: Guide to Molecular Cloning Techniques", Academic Press, Berger, S. L. and A. R. Kimmel eds., 1987.

### 4.10.2 NUTRITIONAL USES

Polynucleotides and polypeptides of the present invention can also be used as nutritional sources or supplements. Such uses include without limitation use as a protein or amino acid supplement, use as a carbon source, use as a nitrogen source and use as a source of carbohydrate. In such cases the polypeptide or polynucleotide of the invention can be added to the feed of a particular organism or can be administered as a separate solid or liquid preparation, such as in the form of powder, pills, solutions, suspensions or capsules. In the case of microorganisms, the polypeptide or polynucleotide of the invention can be added to the medium in or on which the microorganism is cultured.

### 4.10.3 CYTOKINE AND CELL PROLIFERATION/DIFFERENTIATION ACTIVITY

A polypeptide of the present invention may exhibit activity relating to cytokine, cell proliferation (either inducing or inhibiting) or cell differentiation (either inducing or inhibiting) activity or may induce production of other cytokines in certain cell populations. A polynucleotide of the invention can encode a polypeptide exhibiting such attributes. Many protein factors discovered to date, including all known cytokines, have exhibited activity in one or more factor-dependent cell proliferation assays, and hence the assays serve as a convenient

confirmation of cytokine activity. The activity of therapeutic compositions of the present invention is evidenced by any one of a number of routine factor dependent cell proliferation assays for cell lines including, without limitation, 32D, DA2, DA1G, T10, B9, B9/11, BaF3, MC9/G, M+(preB M+), 2E8, RB5, DA1, 123, T1165, HT2, CTLL2, TF-1, Mo7e, CMK, HUVEC, and Caco. Therapeutic compositions of the invention can be used in the following:

Assays for T-cell or thymocyte proliferation include without limitation those described in: Current Protocols in Immunology, Ed by J. E. Coligan, A. M. Kruisbeek, D. H. Margulies, E. M. Shevach, W. Strober, Pub. Greene Publishing Associates and Wiley-Interscience (Chapter 3, *In Vitro* assays for Mouse Lymphocyte Function 3.1-3.19; Chapter 7, Immunologic studies in Humans); Takai et al., J. Immunol. 137:3494-3500, 1986; Bertagnolli et al., J. Immunol. 145:1706-1712, 1990; Bertagnolli et al., Cellular Immunology 133:327-341, 1991; Bertagnolli, et al., I. Immunol. 149:3778-3783, 1992; Bowman et al., I: Immunol. 152:1756-1761, 1994.

Assays for cytokine production and/or proliferation of spleen cells, lymph node cells or thymocytes include, without limitation, those described in: Polyclonal T cell stimulation, Kruisbeek, A. M. and Shevach, E. M. In Current Protocols in Immunology. J. E. e.a. Coligan eds. Vol 1 pp. 3.12.1-3.12.14, John Wiley and Sons, Toronto. 1994; and Measurement of mouse and human interleukin-γ, Schreiber, R. D. In Current Protocols in Immunology. J. E. e.a. Coligan eds. Vol 1 pp. 6.8.1-6.8.8, John Wiley and Sons, Toronto. 1994.

Assays for proliferation and differentiation of hematopoietic and lymphopoietic cells include, without limitation, those described in: Measurement of Human and Murine Interleukin 2 and Interleukin 4, Bottomly, K., Davis, L. S. and Lipsky, P. E. In Current Protocols in Immunology. J. E. e.a. Coligan eds. Vol 1 pp. 6.3.1-6.3.12, John Wiley and Sons, Toronto. 1991; deVries et al., J. Exp. Med. 173:1205-1211, 1991; Moreau et al., Nature 336:690-692, 1988; Greenberger et al., Proc. Natl. Acad. Sci. U.S.A. 80:2931-2938, 1983; Measurement of mouse and human interleukin 6--Nordan, R. In Current Protocols in Immunology. J. E. Coligan eds. Vol 1 pp. 6.6.1-6.6.5, John Wiley and Sons, Toronto. 1991; Smith et al., Proc. Natl. Aced. Sci. U.S.A. 83:1857-1861, 1986; Measurement of human Interleukin 11--Bennett, F., Giannotti, J., Clark, S. C. and Turner, K. J. In Current Protocols in Immunology. J. E. Coligan eds. Vol 1 pp. 6.15.1 John Wiley and Sons, Toronto. 1991; Measurement of mouse and human Interleukin 9--Ciarletta, A., Giannotti, J., Clark, S. C. and Turner, K. J. In Current Protocols in Immunology. J. E. Coligan eds. Vol 1 pp. 6.13.1, John Wiley and Sons, Toronto. 1991.

Assays for T-cell clone responses to antigens (which will identify, among others, proteins that affect APC-T cell interactions as well as direct T-cell effects by measuring proliferation and cytokine production) include, without limitation, those described in: Current Protocols in Immunology, Ed by J. E. Coligan, A. M. Kruisbeek, D. H. Margulies, E. M. Shevach, W Strober,

Pub. Greene Publishing Associates and Wiley-Interscience (Chapter 3, *In Vitro* assays for Mouse Lymphocyte Function; Chapter 6, Cytokines and their cellular receptors; Chapter 7, Immunologic studies in Humans); Weinberger et al., Proc. Natl. Acad. Sci. USA 77:6091-6095, 1980; Weinberger et al., Eur. J. Immun. 11:405-411, 1981; Takai et al., J. Immunol. 137:3494-3500, 1986; Takai et al., J. Immunol. 140:508-512, 1988.

### 4.10.4 STEM CELL GROWTH FACTOR ACTIVITY

A polypeptide of the present invention may exhibit stem cell growth factor activity and be involved in the proliferation, differentiation and survival of pluripotent and totipotent stem cells including primordial germ cells, embryonic stem cells, hematopoietic stem cells and/or germ line stem cells. Administration of the polypeptide of the invention to stem cells *in vivo* or *ex vivo* is expected to maintain and expand cell populations in a totipotential or pluripotential state which would be useful for re-engineering damaged or diseased tissues, transplantation, manufacture of bio-pharmaceuticals and the development of bio-sensors. The ability to produce large quantities of human cells has important working applications for the production of human proteins which currently must be obtained from non-human sources or donors, implantation of cells to treat diseases such as Parkinson's, Alzheimer's and other neurodegenerative diseases; tissues for grafting such as bone marrow, skin, cartilage, tendons, bone, muscle (including cardiac muscle), blood vessels, cornea, neural cells, gastrointestinal cells and others; and organs for transplantation such as kidney, liver, pancreas (including islet cells), heart and lung.

It is contemplated that multiple different exogenous growth factors and/or cytokines may be administered in combination with the polypeptide of the invention to achieve the desired effect, including any of the growth factors listed herein, other stem cell maintenance factors, and specifically including stem cell factor (SCF), leukemia inhibitory factor (LIF), Flt-3 ligand (Flt-3L), any of the interleukins, recombinant soluble IL-6 receptor fused to IL-6, macrophage inflammatory protein 1-alpha (MIP-1-alpha), G-CSF, GM-CSF, thrombopoietin (TPO), platelet factor 4 (PF-4), platelet-derived growth factor (PDGF), neural growth factors and basic fibroblast growth factor (bFGF).

Since totipotent stem cells can give rise to virtually any mature cell type, expansion of these cells in culture will facilitate the production of large quantities of mature cells. Techniques for culturing stem cells are known in the art and administration of polypeptides of the invention, optionally with other growth factors and/or cytokines, is expected to enhance the survival and proliferation of the stem cell populations. This can be accomplished by direct administration of the polypeptide of the invention to the culture medium. Alternatively, stroma cells transfected with a polynucleotide that encodes for the polypeptide of the invention can be used as a feeder

layer for the stem cell populations in culture or in vivo. Stromal support cells for feeder layers may include embryonic bone marrow fibroblasts, bone marrow stromal cells, fetal liver cells, or cultured embryonic fibroblasts (see U.S. Patent No. 5,690,926).

Stem cells themselves can be transfected with a polynucleotide of the invention to induce autocrine expression of the polypeptide of the invention. This will allow for generation of undifferentiated totipotential/pluripotential stem cell lines that are useful as is or that can then be differentiated into the desired mature cell types. These stable cell lines can also serve as a source of undifferentiated totipotential/pluripotential mRNA to create cDNA libraries and templates for polymerase chain reaction experiments. These studies would allow for the isolation and identification of differentially expressed genes in stem cell populations that regulate stem cell proliferation and/or maintenance.

Expansion and maintenance of totipotent stem cell populations will be useful in the treatment of many pathological conditions. For example, polypeptides of the present invention may be used to manipulate stem cells in culture to give rise to neuroepithelial cells that can be used to augment or replace cells damaged by illness, autoimmune disease, accidental damage or genetic disorders. The polypeptide of the invention may be useful for inducing the proliferation of neural cells and for the regeneration of nerve and brain tissue, *i.e.* for the treatment of central and peripheral nervous system diseases and neuropathies, as well as mechanical and traumatic disorders which involve degeneration, death or trauma to neural cells or nerve tissue. In addition, the expanded stem cell populations can also be genetically altered for gene therapy purposes and to decrease host rejection of replacement tissues after grafting or implantation.

Expression of the polypeptide of the invention and its effect on stem cells can also be manipulated to achieve controlled differentiation of the stem cells into more differentiated cell types. A broadly applicable method of obtaining pure populations of a specific differentiated cell type from undifferentiated stem cell populations involves the use of a cell-type specific promoter driving a selectable marker. The selectable marker allows only cells of the desired type to survive. For example, stem cells can be induced to differentiate into cardiomyocytes (Wobus et al., Differentiation, 48: 173-182, (1991); Klug et al., J. Clin. Invest., 98(1): 216-224, (1998)) or skeletal muscle cells (Browder, L. W. In: *Principles of Tissue Engineering eds.* Lanza et al., Academic Press (1997)). Alternatively, directed differentiation of stem cells can be accomplished by culturing the stem cells in the presence of a differentiation factor such as retinoic acid and an antagonist of the polypeptide of the invention which would inhibit the effects of endogenous stem cell factor activity and allow differentiation to proceed.

*In vitro* cultures of stem cells can be used to determine if the polypeptide of the invention exhibits stem cell growth factor activity. Stem cells are isolated from any one of various cell

sources (including hematopoietic stem cells and embryonic stem cells) and cultured on a feeder layer, as described by Thompson et al. Proc. Natl. Acad. Sci, U.S.A., 92: 7844-7848 (1995), in the presence of the polypeptide of the invention alone or in combination with other growth factors or cytokines. The ability of the polypeptide of the invention to induce stem cells proliferation is determined by colony formation on semi-solid support *e.g.* as described by Bernstein et al., Blood, 77: 2316-2321 (1991).

### 4.10.5 HEMATOPOIESIS REGULATING ACTIVITY

A polypeptide of the present invention may be involved in regulation of hematopoiesis and, consequently, in the treatment of myeloid or lymphoid cell disorders. Even marginal biological activity in support of colony forming cells or of factor-dependent cell lines indicates involvement in regulating hematopoiesis, *e.g.* in supporting the growth and proliferation of erythroid progenitor cells alone or in combination with other cytokines, thereby indicating utility, for example, in treating various anemias or for use in conjunction with irradiation/chemotherapy to stimulate the production of erythroid precursors and/or erythroid cells; in supporting the growth and proliferation of myeloid cells such as granulocytes and monocytes/macrophages (*i.e.*, traditional CSF activity) useful, for example, in conjunction with chemotherapy to prevent or treat consequent myelo-suppression; in supporting the growth and proliferation of megakaryocytes and consequently of platelets thereby allowing prevention or treatment of various platelet disorders such as thrombocytopenia, and generally for use in place of or complimentary to platelet transfusions; and/or in supporting the growth and proliferation of hematopoietic stem cells which are capable of maturing to any and all of the above-mentioned hematopoietic cells and therefore find therapeutic utility in various stem cell disorders (such as those usually treated with transplantation, including, without limitation, aplastic anemia and paroxysmal nocturnal hemoglobinuria), as well as in repopulating the stem cell compartment post irradiation/chemotherapy, either *in-vivo* or *ex-vivo* (*i.e.*, in conjunction with bone marrow transplantation or with peripheral progenitor cell transplantation (homologous or heterologous)) as normal cells or genetically manipulated for gene therapy.

Therapeutic compositions of the invention can be used in the following:

Suitable assays for proliferation and differentiation of various hematopoietic lines are cited above.

Assays for embryonic stem cell differentiation (which will identify, among others, proteins that influence embryonic differentiation hematopoiesis) include, without limitation, those described in: Johansson et al. Cellular Biology 15:141-151, 1995; Keller et al., Molecular and Cellular Biology 13:473-486, 1993; McClanahan et al., Blood 81:2903-2915, 1993.

Assays for stem cell survival and differentiation (which will identify, among others, proteins that regulate lympho-hematopoiesis) include, without limitation, those described in: Methylcellulose colony forming assays, Freshney, M. G. In Culture of Hematopoietic Cells. R. I. Freshney, et al. eds. Vol pp. 265-268, Wiley-Liss, Inc., New York, N.Y. 1994; Hirayama et al., Proc. Natl. Acad. Sci. USA 89:5907-5911, 1992; Primitive hematopoietic colony forming cells with high proliferative potential, McNiece, I. K. and Briddell, R. A. In Culture of Hematopoietic Cells. R. I. Freshney, et al. eds. Vol pp. 23-39, Wiley-Liss, Inc., New York, N.Y. 1994; Neben et al., Experimental Hematology 22:353-359, 1994; Cobblestone area forming cell assay, Ploemacher, R. E. In Culture of Hematopoietic Cells. R. I. Freshney, et al. eds. Vol pp. 1-21, Wiley-Liss, Inc., New York, N.Y. 1994; Long term bone marrow cultures in the presence of stromal cells, Spooncer, E., Dexter, M. and Allen, T. In Culture of Hematopoietic Cells. R. I. Freshney, et al. eds. Vol pp. 163-179, Wiley-Liss, Inc., New York, N.Y. 1994; Long term culture initiating cell assay, Sutherland, H. J. In Culture of Hematopoietic Cells. R. I. Freshney, et al. eds. Vol pp. 139-162, Wiley-Liss, Inc., New York, N.Y. 1994.

### 4.10.6 TISSUE GROWTH ACTIVITY

A polypeptide of the present invention also may be involved in bone, cartilage, tendon, ligament and/or nerve tissue growth or regeneration, as well as in wound healing and tissue repair and replacement, and in healing of burns, incisions and ulcers.

A polypeptide of the present invention which induces cartilage and/or bone growth in circumstances where bone is not normally formed, has application in the healing of bone fractures and cartilage damage or defects in humans and other animals. Compositions of a polypeptide, antibody, binding partner, or other modulator of the invention may have prophylactic use in closed as well as open fracture reduction and also in the improved fixation of artificial joints. De novo bone formation induced by an osteogenic agent contributes to the repair of congenital, trauma induced, or oncologic resection induced craniofacial defects, and also is useful in cosmetic plastic surgery.

A polypeptide of this invention may also be involved in attracting bone-forming cells, stimulating growth of bone-forming cells, or inducing differentiation of progenitors of bone-forming cells. Treatment of osteoporosis, osteoarthritis, bone degenerative disorders, or periodontal disease, such as through stimulation of bone and/or cartilage repair or by blocking inflammation or processes of tissue destruction (collagenase activity, osteoclast activity, etc.) mediated by inflammatory processes may also be possible using the composition of the invention.

Another category of tissue regeneration activity that may involve the polypeptide of the present invention is tendon/ligament formation. Induction of tendon/ligament-like tissue or other tissue formation in circumstances where such tissue is not normally formed, has application in the healing of tendon or ligament tears, deformities and other tendon or ligament defects in humans and other animals. Such a preparation employing a tendon/ligament-like tissue inducing protein may have prophylactic use in preventing damage to tendon or ligament tissue, as well as use in the improved fixation of tendon or ligament to bone or other tissues, and in repairing defects to tendon or ligament tissue. De novo tendon/ligament-like tissue formation induced by a composition of the present invention contributes to the repair of congenital, trauma induced, or other tendon or ligament defects of other origin, and is also useful in cosmetic plastic surgery for attachment or repair of tendons or ligaments. The compositions of the present invention may provide environment to attract tendon- or ligament-forming cells, stimulate growth of tendon- or ligament-forming cells, induce differentiation of progenitors of tendon- or ligament-forming cells, or induce growth of tendon/ligament cells or progenitors *ex vivo* for return *in vivo* to effect tissue repair. The compositions of the invention may also be useful in the treatment of tendinitis, carpal tunnel syndrome and other tendon or ligament defects. The compositions may also include an appropriate matrix and/or sequestering agent as a carrier as is well known in the art.

The compositions of the present invention may also be useful for proliferation of neural cells and for regeneration of nerve and brain tissue, *i.e.* for the treatment of central and peripheral nervous system diseases and neuropathies, as well as mechanical and traumatic disorders, which involve degeneration, death or trauma to neural cells or nerve tissue. More specifically, a composition may be used in the treatment of diseases of the peripheral nervous system, such as peripheral nerve injuries, peripheral neuropathy and localized neuropathies, and central nervous system diseases, such as Alzheimer's, Parkinson's disease, Huntington's disease, amyotrophic lateral sclerosis, and Shy-Drager syndrome. Further conditions which may be treated in accordance with the present invention include mechanical and traumatic disorders, such as spinal cord disorders, head trauma and cerebrovascular diseases such as stroke. Peripheral neuropathies resulting from chemotherapy or other medical therapies may also be treatable using a composition of the invention.

Compositions of the invention may also be useful to promote better or faster closure of non-healing wounds, including without limitation pressure ulcers, ulcers associated with vascular insufficiency, surgical and traumatic wounds, and the like.

Compositions of the present invention may also be involved in the generation or regeneration of other tissues, such as organs (including, for example, pancreas, liver, intestine,

kidney, skin, endothelium), muscle (smooth, skeletal or cardiac) and vascular (including vascular endothelium) tissue, or for promoting the growth of cells comprising such tissues. Part of the desired effects may be by inhibition or modulation of fibrotic scarring may allow normal tissue to regenerate. A polypeptide of the present invention may also exhibit angiogenic activity.

A composition of the present invention may also be useful for gut protection or regeneration and treatment of lung or liver fibrosis, reperfusion injury in various tissues, and conditions resulting from systemic cytokine damage.

A composition of the present invention may also be useful for promoting or inhibiting differentiation of tissues described above from precursor tissues or cells; or for inhibiting the growth of tissues described above.

Therapeutic compositions of the invention can be used in the following:

Assays for tissue generation activity include, without limitation, those described in: International Patent Publication No. WO95/16035 (bone, cartilage, tendon); International Patent Publication No. WO95/05846 (nerve, neuronal); International Patent Publication No. WO91/07491 (skin, endothelium).

Assays for wound healing activity include, without limitation, those described in: Winter, Epidermal Wound Healing, pps. 71-112 (Maibach, H. I. and Rovee, D. T., eds.), Year Book Medical Publishers, Inc., Chicago, as modified by Eaglstein and Mertz, J. Invest. Dermatol 71:382-84 (1978).

### 4.10.7 IMMUNE STIMULATING OR SUPPRESSING ACTIVITY

A polypeptide of the present invention may also exhibit immune stimulating or immune suppressing activity, including without limitation the activities for which assays are described herein. A polynucleotide of the invention can encode a polypeptide exhibiting such activities. A protein may be useful in the treatment of various immune deficiencies and disorders (including severe combined immunodeficiency (SCID)), *e.g.*, in regulating (up or down) growth and proliferation of T and/or B lymphocytes, as well as effecting the cytolytic activity of NK cells and other cell populations. These immune deficiencies may be genetic or be caused by viral (*e.g.*, HIV) as well as bacterial or fungal infections, or may result from autoimmune disorders. More specifically, infectious diseases causes by viral, bacterial, fungal or other infection may be treatable using a protein of the present invention, including infections by HIV, hepatitis viruses, herpes viruses, mycobacteria, Leishmania spp., malaria spp. and various fungal infections such as candidiasis. Of course, in this regard, proteins of the present invention may also be useful where a boost to the immune system generally may be desirable, *i.e.*, in the treatment of cancer.

Autoimmune disorders which may be treated using a protein of the present invention include, for example, connective tissue disease, multiple sclerosis, systemic lupus erythematosus, rheumatoid arthritis, autoimmune pulmonary inflammation, Guillain-Barre syndrome, autoimmune thyroiditis, insulin dependent diabetes mellitis, myasthenia gravis, graft-versus-host disease and autoimmune inflammatory eye disease. Such a protein (or antagonists thereof, including antibodies) of the present invention may also to be useful in the treatment of allergic reactions and conditions (*e.g.*, anaphylaxis, serum sickness, drug reactions, food allergies, insect venom allergies, mastocytosis, allergic rhinitis, hypersensitivity pneumonitis, urticaria, angioedema, eczema, atopic dermatitis, allergic contact dermatitis, erythema multiforme, Stevens-Johnson syndrome, allergic conjunctivitis, atopic keratoconjunctivitis, venereal keratoconjunctivitis, giant papillary conjunctivitis and contact allergies), such as asthma (particularly allergic asthma) or other respiratory problems. Other conditions, in which immune suppression is desired (including, for example, organ transplantation), may also be treatable using a protein (or antagonists thereof) of the present invention. The therapeutic effects of the polypeptides or antagonists thereof on allergic reactions can be evaluated by in vivo animals models such as the cumulative contact enhancement test (Lastbom et al., Toxicology 125: 59-66, 1998), skin prick test (Hoffmann et al., Allergy 54: 446-54, 1999), guinea pig skin sensitization test (Vohr et al., Arch. Toxocol. 73: 501-9), and murine local lymph node assay (Kimber et al., J. Toxicol. Environ. Health 53: 563-79).

Using the proteins of the invention it may also be possible to modulate immune responses, in a number of ways. Down regulation may be in the form of inhibiting or blocking an immune response already in progress or may involve preventing the induction of an immune response. The functions of activated T cells may be inhibited by suppressing T cell responses or by inducing specific tolerance in T cells, or both. Immunosuppression of T cell responses is generally an active, non-antigen-specific, process which requires continuous exposure of the T cells to the suppressive agent. Tolerance, which involves inducing non-responsiveness or anergy in T cells, is distinguishable from immunosuppression in that it is generally antigen-specific and persists after exposure to the tolerizing agent has ceased. Operationally, tolerance can be demonstrated by the lack of a T cell response upon reexposure to specific antigen in the absence of the tolerizing agent.

Down regulating or preventing one or more antigen functions (including without limitation B lymphocyte antigen functions (such as, for example, B7)), *e.g.*, preventing high level lymphokine synthesis by activated T cells, will be useful in situations of tissue, skin and organ transplantation and in graft-versus-host disease (GVHD). For example, blockage of T cell function should result in reduced tissue destruction in tissue transplantation. Typically, in tissue

transplants, rejection of the transplant is initiated through its recognition as foreign by T cells, followed by an immune reaction that destroys the transplant. The administration of a therapeutic composition of the invention may prevent cytokine synthesis by immune cells, such as T cells, and thus acts as an immunosuppressant. Moreover, a lack of costimulation may also be sufficient to anergize the T cells, thereby inducing tolerance in a subject. Induction of long-term tolerance by B lymphocyte antigen-blocking reagents may avoid the necessity of repeated administration of these blocking reagents. To achieve sufficient immunosuppression or tolerance in a subject, it may also be necessary to block the function of a combination of B lymphocyte antigens.

The efficacy of particular therapeutic compositions in preventing organ transplant rejection or GVHD can be assessed using animal models that are predictive of efficacy in humans. Examples of appropriate systems which can be used include allogeneic cardiac grafts in rats and xenogeneic pancreatic islet cell grafts in mice, both of which have been used to examine the immunosuppressive effects of CTLA4Ig fusion proteins in vivo as described in Lenschow et al., Science 257:789-792 (1992) and Turka et al., Proc. Natl. Acad. Sci USA, 89:11102-11105 (1992). In addition, murine models of GVHD (see Paul ed., Fundamental Immunology, Raven Press, New York, 1989, pp. 846-847) can be used to determine the effect of therapeutic compositions of the invention on the development of that disease.

Blocking antigen function may also be therapeutically useful for treating autoimmune diseases. Many autoimmune disorders are the result of inappropriate activation of T cells that are reactive against self tissue and which promote the production of cytokines and autoantibodies involved in the pathology of the diseases. Preventing the activation of autoreactive T cells may reduce or eliminate disease symptoms. Administration of reagents which block stimulation of T cells can be used to inhibit T cell activation and prevent production of autoantibodies or T cell-derived cytokines which may be involved in the disease process. Additionally, blocking reagents may induce antigen-specific tolerance of autoreactive T cells which could lead to long-term relief from the disease. The efficacy of blocking reagents in preventing or alleviating autoimmune disorders can be determined using a number of well-characterized animal models of human autoimmune diseases. Examples include murine experimental autoimmune encephalitis, systemic lupus erythmatosis in MRL/lpr/lpr mice or NZB hybrid mice, murine autoimmune collagen arthritis, diabetes mellitus in NOD mice and BB rats, and murine experimental myasthenia gravis (see Paul ed., Fundamental Immunology, Raven Press, New York, 1989, pp. 840-856).

Upregulation of an antigen function (*e.g.*, a B lymphocyte antigen function), as a means of up regulating immune responses, may also be useful in therapy. Upregulation of immune responses may be in the form of enhancing an existing immune response or eliciting an initial

immune response. For example, enhancing an immune response may be useful in cases of viral infection, including systemic viral diseases such as influenza, the common cold, and encephalitis.

Alternatively, anti-viral immune responses may be enhanced in an infected patient by removing T cells from the patient, costimulating the T cells in vitro with viral antigen-pulsed APCs either expressing a peptide of the present invention or together with a stimulatory form of a soluble peptide of the present invention and reintroducing the in vitro activated T cells into the patient. Another method of enhancing anti-viral immune responses would be to isolate infected cells from a patient, transfect them with a nucleic acid encoding a protein of the present invention as described herein such that the cells express all or a portion of the protein on their surface, and reintroduce the transfected cells into the patient. The infected cells would now be capable of delivering a costimulatory signal to, and thereby activate, T cells in vivo.

A polypeptide of the present invention may provide the necessary stimulation signal to T cells to induce a T cell mediated immune response against the transfected tumor cells. In addition, tumor cells which lack MHC class I or MHC class II molecules, or which fail to reexpress sufficient mounts of MHC class I or MHC class II molecules, can be transfected with nucleic acid encoding all or a portion of (*e.g.*, a cytoplasmic-domain truncated portion) of an MHC class I alpha chain protein and $\beta_2$ microglobulin protein or an MHC class II alpha chain protein and an MHC class II beta chain protein to thereby express MHC class I or MHC class II proteins on the cell surface. Expression of the appropriate class I or class II MHC in conjunction with a peptide having the activity of a B lymphocyte antigen (*e.g.*, B7-1, B7-2, B7-3) induces a T cell mediated immune response against the transfected tumor cell. Optionally, a gene encoding an antisense construct which blocks expression of an MHC class II associated protein, such as the invariant chain, can also be cotransfected with a DNA encoding a peptide having the activity of a B lymphocyte antigen to promote presentation of tumor associated antigens and induce tumor specific immunity. Thus, the induction of a T cell mediated immune response in a human subject may be sufficient to overcome tumor-specific tolerance in the subject.

The activity of a protein of the invention may, among other means, be measured by the following methods:

Suitable assays for thymocyte or splenocyte cytotoxicity include, without limitation, those described in: Current Protocols in Immunology, Ed by J. E. Coligan, A. M. Kruisbeek, D. H. Margulies, E. M. Shevach, W. Strober, Pub. Greene Publishing Associates and Wiley-Interscience (Chapter 3, In Vitro assays for Mouse Lymphocyte Function 3.1-3.19; Chapter 7, Immunologic studies in Humans); Herrmann et al., Proc. Natl. Acad. Sci. USA 78:2488-2492, 1981; Herrmann et al., J. Immunol. 128:1968-1974, 1982; Handa et al., J.

Immunol. 135:1564-1572, 1985; Takai et al., I. Immunol. 137:3494-3500, 1986; Takai et al., J. Immunol. 140:508-512, 1988; Bowman et al., J. Virology 61:1992-1998; Bertagnolli et al., Cellular Immunology 133:327-341, 1991; Brown et al., J. Immunol. 153:3079-3092, 1994.

Assays for T-cell-dependent immunoglobulin responses and isotype switching (which will identify, among others, proteins that modulate T-cell dependent antibody responses and that affect Th1/Th2 profiles) include, without limitation, those described in: Maliszewski, J. Immunol. 144:3028-3033, 1990; and Assays for B cell function: In vitro antibody production, Mond, J. J. and Brunswick, M. In Current Protocols in Immunology. J. E. e.a. Coligan eds. Vol 1 pp. 3.8.1-3.8.16, John Wiley and Sons, Toronto. 1994.

Mixed lymphocyte reaction (MLR) assays (which will identify, among others, proteins that generate predominantly Th1 and CTL responses) include, without limitation, those described in: Current Protocols in Immunology, Ed by J. E. Coligan, A. M. Kruisbeek, D. H. Margulies, E. M. Shevach, W. Strober, Pub. Greene Publishing Associates and Wiley-Interscience (Chapter 3, In Vitro assays for Mouse Lymphocyte Function 3.1-3.19; Chapter 7, Immunologic studies in Humans); Takai et al., J. Immunol. 137:3494-3500, 1986; Takai et al., J. Immunol. 140:508-512, 1988; Bertagnolli et al., J. Immunol. 149:3778-3783, 1992.

Dendritic cell-dependent assays (which will identify, among others, proteins expressed by dendritic cells that activate naive T-cells) include, without limitation, those described in: Guery et al., J. Immunol. 134:536-544, 1995; Inaba et al., Journal of Experimental Medicine 173:549-559, 1991; Macatonia et al., Journal of Immunology 154:5071-5079, 1995; Porgador et al., Journal of Experimental Medicine 182:255-260, 1995; Nair et al., Journal of Virology 67:4062-4069, 1993; Huang et al., Science 264:961-965, 1994; Macatonia et al., Journal of Experimental Medicine 169:1255-1264, 1989; Bhardwaj et al., Journal of Clinical Investigation 94:797-807, 1994; and Inaba et al., Journal of Experimental Medicine 172:631-640, 1990.

Assays for lymphocyte survival/apoptosis (which will identify, among others, proteins that prevent apoptosis after superantigen induction and proteins that regulate lymphocyte homeostasis) include, without limitation, those described in: Darzynkiewicz et al., Cytometry 13:795-808, 1992; Gorczyca et al., Leukemia 7:659-670, 1993; Gorczyca et al., Cancer Research 53:1945-1951, 1993; Itoh et al., Cell 66:233-243, 1991; Zacharchuk, Journal of Immunology 145:4037-4045, 1990; Zamai et al., Cytometry 14:891-897, 1993; Gorczyca et al., International Journal of Oncology 1:639-648, 1992.

Assays for proteins that influence early steps of T-cell commitment and development include, without limitation, those described in: Antica et al., Blood 84:111-117, 1994; Fine et al., Cellular Immunology 155:111-122, 1994; Galy et al., Blood 85:2770-2778, 1995; Toki et al., Proc. Nat. Acad Sci. USA 88:7548-7551, 1991.

### 4.10.8 ACTIVIN/INHIBIN ACTIVITY

A polypeptide of the present invention may also exhibit activin- or inhibin-related activities. A polynucleotide of the invention may encode a polypeptide exhibiting such characteristics. Inhibins are characterized by their ability to inhibit the release of follicle stimulating hormone (FSH), while activins and are characterized by their ability to stimulate the release of follicle stimulating hormone (FSH). Thus, a polypeptide of the present invention, alone or in heterodimers with a member of the inhibin family, may be useful as a contraceptive based on the ability of inhibins to decrease fertility in female mammals and decrease spermatogenesis in male mammals. Administration of sufficient amounts of other inhibins can induce infertility in these mammals. Alternatively, the polypeptide of the invention, as a homodimer or as a heterodimer with other protein subunits of the inhibin group, may be useful as a fertility inducing therapeutic, based upon the ability of activin molecules in stimulating FSH release from cells of the anterior pituitary. See, for example, U.S. Pat. No. 4,798,885. A polypeptide of the invention may also be useful for advancement of the onset of fertility in sexually immature mammals, so as to increase the lifetime reproductive performance of domestic animals such as, but not limited to, cows, sheep and pigs.

The activity of a polypeptide of the invention may, among other means, be measured by the following methods.

Assays for activin/inhibin activity include, without limitation, those described in: Vale et al., Endocrinology 91:562-572, 1972; Ling et al., Nature 321:779-782, 1986; Vale et al., Nature 321:776-779, 1986; Mason et al., Nature 318:659-663, 1985; Forage et al., Proc. Natl. Acad. Sci. USA 83:3091-3095, 1986.

### 4.10.9 CHEMOTACTIC/CHEMOKINETIC ACTIVITY

A polypeptide of the present invention may be involved in chemotactic or chemokinetic activity for mammalian cells, including, for example, monocytes, fibroblasts, neutrophils, T-cells, mast cells, eosinophils, epithelial and/or endothelial cells. A polynucleotide of the invention can encode a polypeptide exhibiting such attributes. Chemotactic and chemokinetic receptor activation can be used to mobilize or attract a desired cell population to a desired site of action. Chemotactic or chemokinetic compositions (*e.g.* proteins, antibodies, binding partners, or modulators of the invention) provide particular advantages in treatment of wounds and other trauma to tissues, as well as in treatment of localized infections. For example, attraction of lymphocytes, monocytes or neutrophils to tumors or sites of infection may result in improved immune responses against the tumor or infecting agent.

A protein or peptide has chemotactic activity for a particular cell population if it can stimulate, directly or indirectly, the directed orientation or movement of such cell population. Preferably, the protein or peptide has the ability to directly stimulate directed movement of cells. Whether a particular protein has chemotactic activity for a population of cells can be readily determined by employing such protein or peptide in any known assay for cell chemotaxis.

Therapeutic compositions of the invention can be used in the following:

Assays for chemotactic activity (which will identify proteins that induce or prevent chemotaxis) consist of assays that measure the ability of a protein to induce the migration of cells across a membrane as well as the ability of a protein to induce the adhesion of one cell population to another cell population. Suitable assays for movement and adhesion include, without limitation, those described in: Current Protocols in Immunology, Ed by J. E. Coligan, A. M. Kruisbeek, D. H. Marguiles, E. M. Shevach, W. Strober, Pub. Greene Publishing Associates and Wiley-Interscience (Chapter 6.12, Measurement of alpha and beta Chemokines 6.12.1-6.12.28; Taub et al. J. Clin. Invest. 95:1370-1376, 1995; Lind et al. APMIS 103:140-146, 1995; Muller et al Eur. J. Immunol. 25:1744-1748; Gruber et al. J. of Immunol. 152:5860-5867, 1994; Johnston et al. J. of Immunol. 153:1762-1768, 1994.

### 4.10.10 HEMOSTATIC AND THROMBOLYTIC ACTIVITY

A polypeptide of the invention may also be involved in hemostatis or thrombolysis or thrombosis. A polynucleotide of the invention can encode a polypeptide exhibiting such attributes. Compositions may be useful in treatment of various coagulation disorders (including hereditary disorders, such as hemophilias) or to enhance coagulation and other hemostatic events in treating wounds resulting from trauma, surgery or other causes. A composition of the invention may also be useful for dissolving or inhibiting formation of thromboses and for treatment and prevention of conditions resulting therefrom (such as, for example, infarction of cardiac and central nervous system vessels (*e.g.*, stroke).

Therapeutic compositions of the invention can be used in the following:

Assay for hemostatic and thrombolytic activity include, without limitation, those described in: Linet et al., J. Clin. Pharmacol. 26:131-140, 1986; Burdick et al., Thrombosis Res. 45:413-419, 1987; Humphrey et al., Fibrinolysis 5:71-79 (1991); Schaub, Prostaglandins 35:467-474, 1988.

### 4.10.11 CANCER DIAGNOSIS AND THERAPY

Polypeptides of the invention may be involved in cancer cell generation, proliferation or metastasis. Detection of the presence or amount of polynucleotides or polypeptides of the

invention may be useful for the diagnosis and/or prognosis of one or more types of cancer. For example, the presence or increased expression of a polynucleotide/polypeptide of the invention may indicate a hereditary risk of cancer, a precancerous condition, or an ongoing malignancy. Conversely, a defect in the gene or absence of the polypeptide may be associated with a cancer condition. Identification of single nucleotide polymorphisms associated with cancer or a predisposition to cancer may also be useful for diagnosis or prognosis.

Cancer treatments promote tumor regression by inhibiting tumor cell proliferation, inhibiting angiogenesis (growth of new blood vessels that is necessary to support tumor growth) and/or prohibiting metastasis by reducing tumor cell motility or invasiveness. Therapeutic compositions of the invention may be effective in adult and pediatric oncology including in solid phase tumors/malignancies, locally advanced tumors, human soft tissue sarcomas, metastatic cancer, including lymphatic metastases, blood cell malignancies including multiple myeloma, acute and chronic leukemias, and lymphomas, head and neck cancers including mouth cancer, larynx cancer and thyroid cancer, lung cancers including small cell carcinoma and non-small cell cancers, breast cancers including small cell carcinoma and ductal carcinoma, gastrointestinal cancers including esophageal cancer, stomach cancer, colon cancer, colorectal cancer and polyps associated with colorectal neoplasia, pancreatic cancers, liver cancer, urologic cancers including bladder cancer and prostate cancer, malignancies of the female genital tract including ovarian carcinoma, uterine (including endometrial) cancers, and solid tumor in the ovarian follicle, kidney cancers including renal cell carcinoma, brain cancers including intrinsic brain tumors, neuroblastoma, astrocytic brain tumors, gliomas, metastatic tumor cell invasion in the central nervous system, bone cancers including osteomas, skin cancers including malignant melanoma, tumor progression of human skin keratinocytes, squamous cell carcinoma, basal cell carcinoma, hemangiopericytoma and Karposi's sarcoma.

Polypeptides, polynucleotides, or modulators of polypeptides of the invention (including inhibitors and stimulators of the biological activity of the polypeptide of the invention) may be administered to treat cancer. Therapeutic compositions can be administered in therapeutically effective dosages alone or in combination with adjuvant cancer therapy such as surgery, chemotherapy, radiotherapy, thermotherapy, and laser therapy, and may provide a beneficial effect, *e.g.* reducing tumor size, slowing rate of tumor growth, inhibiting metastasis, or otherwise improving overall clinical condition, without necessarily eradicating the cancer.

The composition can also be administered in therapeutically effective amounts as a portion of an anti-cancer cocktail. An anti-cancer cocktail is a mixture of the polypeptide or modulator of the invention with one or more anti-cancer drugs in addition to a pharmaceutically acceptable carrier for delivery. The use of anti-cancer cocktails as a cancer treatment is routine.

Anti-cancer drugs that are well known in the art and can be used as a treatment in combination with the polypeptide or modulator of the invention include: Actinomycin D, Aminoglütethimide, Asparaginase, Bleomycin, Busulfan, Carboplatin, Carmustine, Chlorambucil, Cisplatin (cis-DDP), Cyclophosphamide, Cytarabine HCl (Cytosine arabinoside), Dacarbazine, Dactinomycin, Daunorubicin HCl, Doxorubicin HCl, Estramustine phosphate sodium, Etoposide (V16-213), Floxuridine, 5-Fluorouracil (5-Fu), Flutamide, Hydroxyurea (hydroxycarbamide), Ifosfamide, Interferon Alpha-2a, Interferon Alpha-2b, Leuprolide acetate (LHRH-releasing factor analog), Lomustine, Mechlorethamine HCl (nitrogen mustard), Melphalan, Mercaptopurine, Mesna, Methotrexate (MTX), Mitomycin, Mitoxantrone HCl, Octreotide, Plicamycin, Procarbazine HCl, Streptozocin, Tamoxifen citrate, Thioguanine, Thiotepa, Vinblastine sulfate, Vincristine sulfate, Amsacrine, Azacitidine, Hexamethylmelamine, Interleukin-2, Mitoguazone, Pentostatin, Semustine, Teniposide, and Vindesine sulfate.

In addition, therapeutic compositions of the invention may be used for prophylactic treatment of cancer. There are hereditary conditions and/or environmental situations (*e.g.* exposure to carcinogens) known in the art that predispose an individual to developing cancers. Under these circumstances, it may be beneficial to treat these individuals with therapeutically effective doses of the polypeptide of the invention to reduce the risk of developing cancers.

*In vitro* models can be used to determine the effective doses of the polypeptide of the invention as a potential cancer treatment. These *in vitro* models include proliferation assays of cultured tumor cells, growth of cultured tumor cells in soft agar (see Freshney, (1987) Culture of Animal Cells: A Manual of Basic Technique, Wily-Liss, New York, NY Ch 18 and Ch 21), tumor systems in nude mice as described in Giovanella et al., J. Natl. Can. Inst., 52: 921-30 (1974), mobility and invasive potential of tumor cells in Boyden Chamber assays as described in Pilkington et al., Anticancer Res., 17: 4107-9 (1997), and angiogenesis assays such as induction of vascularization of the chick chorioallantoic membrane or induction of vascular endothelial cell migration as described in Ribatta et al., Intl. J. Dev. Biol., 40: 1189-97 (1999) and Li et al., Clin. Exp. Metastasis, 17:423-9 (1999), respectively. Suitable tumor cells lines are available, *e.g.* from American Type Tissue Culture Collection catalogs.

### 4.10.12 RECEPTOR/LIGAND ACTIVITY

A polypeptide of the present invention may also demonstrate activity as receptor, receptor ligand or inhibitor or agonist of receptor/ligand interactions. A polynucleotide of the invention can encode a polypeptide exhibiting such characteristics. Examples of such receptors and ligands include, without limitation, cytokine receptors and their ligands, receptor kinases and their ligands, receptor phosphatases and their ligands, receptors involved in cell-cell interactions

and their ligands (including without limitation, cellular adhesion molecules (such as selectins, integrins and their ligands) and receptor/ligand pairs involved in antigen presentation, antigen recognition and development of cellular and humoral immune responses. Receptors and ligands are also useful for screening of potential peptide or small molecule inhibitors of the relevant receptor/ligand interaction. A protein of the present invention (including, without limitation, fragments of receptors and ligands) may themselves be useful as inhibitors of receptor/ligand interactions.

The activity of a polypeptide of the invention may, among other means, be measured by the following methods:

Suitable assays for receptor-ligand activity include without limitation those described in: Current Protocols in Immunology, Ed by J. E. Coligan, A. M. Kruisbeek, D. H. Margulies, E. M. Shevach, W. Strober, Pub. Greene Publishing Associates and Wiley- Interscience (Chapter 7.28, Measurement of Cellular Adhesion under static conditions 7.28.1- 7.28.22), Takai et al., Proc. Natl. Acad. Sci. USA 84:6864-6868, 1987; Bierer et al., J. Exp. Med. 168:1145-1156, 1988; Rosenstein et al., J. Exp. Med. 169:149-160 1989; Stoltenborg et al., J. Immunol. Methods 175:59-68, 1994; Stitt et al., Cell 80:661-670, 1995.

By way of example, the polypeptides of the invention may be used as a receptor for a ligand(s) thereby transmitting the biological activity of that ligand(s). Ligands may be identified through binding assays, affinity chromatography, dihybrid screening assays, BIAcore assays, gel overlay assays, or other methods known in the art.

Studies characterizing drugs or proteins as agonist or antagonist or partial agonists or a partial antagonist require the use of other proteins as competing ligands. The polypeptides of the present invention or ligand(s) thereof may be labeled by being coupled to radioisotopes, colorimetric molecules or a toxin molecules by conventional methods. ("Guide to Protein Purification" Murray P. Deutscher (ed) Methods in Enzymology Vol. 182 (1990) Academic Press, Inc. San Diego). Examples of radioisotopes include, but are not limited to, tritium and carbon-14 . Examples of colorimetric molecules include, but are not limited to, fluorescent molecules such as fluorescamine, or rhodamine or other colorimetric molecules. Examples of toxins include, but are not limited, to ricin.

### 4.10.13 DRUG SCREENING

This invention is particularly useful for screening chemical compounds by using the novel polypeptides or binding fragments thereof in any of a variety of drug screening techniques. The polypeptides or fragments employed in such a test may either be free in solution, affixed to a solid support, borne on a cell surface or located intracellularly. One method of drug screening

utilizes eukaryotic or prokaryotic host cells which are stably transformed with recombinant nucleic acids expressing the polypeptide or a fragment thereof. Drugs are screened against such transformed cells in competitive binding assays. Such cells, either in viable or fixed form, can be used for standard binding assays. One may measure, for example, the formation of complexes between polypeptides of the invention or fragments and the agent being tested or examine the diminution in complex formation between the novel polypeptides and an appropriate cell line, which are well known in the art.

Sources for test compounds that may be screened for ability to bind to or modulate (*i.e.*, increase or decrease) the activity of polypeptides of the invention include (1) inorganic and organic chemical libraries, (2) natural product libraries, and (3) combinatorial libraries comprised of either random or mimetic peptides, oligonucleotides or organic molecules.

Chemical libraries may be readily synthesized or purchased from a number of commercial sources, and may include structural analogs of known compounds or compounds that are identified as "hits" or "leads" via natural product screening.

The sources of natural product libraries are microorganisms (including bacteria and fungi), animals, plants or other vegetation, or marine organisms, and libraries of mixtures for screening may be created by: (1) fermentation and extraction of broths from soil, plant or marine microorganisms or (2) extraction of the organisms themselves. Natural product libraries include polyketides, non-ribosomal peptides, and (non-naturally occurring) variants thereof. For a review, see *Science 282*:63-68 (1998).

Combinatorial libraries are composed of large numbers of peptides, oligonucleotides or organic compounds and can be readily prepared by traditional automated synthesis methods, PCR, cloning or proprietary synthetic methods. Of particular interest are peptide and oligonucleotide combinatorial libraries. Still other libraries of interest include peptide, protein, peptidomimetic, multiparallel synthetic collection, recombinatorial, and polypeptide libraries. For a review of combinatorial chemistry and libraries created therefrom, see Myers, *Curr. Opin. Biotechnol.* 8:701-707 (1997). For reviews and examples of peptidomimetic libraries, see Al-Obeidi et al., *Mol. Biotechnol*, 9(3):205-23 (1998); Hruby et al., *Curr Opin Chem Biol*, 1(1):114-19 (1997); Dorner et al., *Bioorg Med Chem*, 4(5):709-15 (1996) (alkylated dipeptides).

Identification of modulators through use of the various libraries described herein permits modification of the candidate "hit" (or "lead") to optimize the capacity of the "hit" to bind a polypeptide of the invention. The molecules identified in the binding assay are then tested for antagonist or agonist activity in *in vivo* tissue culture or animal models that are well known in the art. In brief, the molecules are titrated into a plurality of cell cultures or animals and then tested for either cell/animal death or prolonged survival of the animal/cells.

The binding molecules thus identified may be complexed with toxins, *e.g.*, ricin or cholera, or with other compounds that are toxic to cells such as radioisotopes. The toxin-binding molecule complex is then targeted to a tumor or other cell by the specificity of the binding molecule for a polypeptide of the invention. Alternatively, the binding molecules may be complexed with imaging agents for targeting and imaging purposes.

### 4.10.14 ASSAY FOR RECEPTOR ACTIVITY

The invention also provides methods to detect specific binding of a polypeptide *e.g.* a ligand or a receptor. The art provides numerous assays particularly useful for identifying previously unknown binding partners for receptor polypeptides of the invention. For example, expression cloning using mammalian or bacterial cells, or dihybrid screening assays can be used to identify polynucleotides encoding binding partners. As another example, affinity chromatography with the appropriate immobilized polypeptide of the invention can be used to isolate polypeptides that recognize and bind polypeptides of the invention. There are a number of different libraries used for the identification of compounds, and in particular small molecules, that modulate (*i.e.,* increase or decrease) biological activity of a polypeptide of the invention. Ligands for receptor polypeptides of the invention can also be identified by adding exogenous ligands, or cocktails of ligands to two cells populations that are genetically identical except for the expression of the receptor of the invention: one cell population expresses the receptor of the invention whereas the other does not. The response of the two cell populations to the addition of ligands(s) are then compared. Alternatively, an expression library can be co-expressed with the polypeptide of the invention in cells and assayed for an autocrine response to identify potential ligand(s). As still another example, BIAcore assays, gel overlay assays, or other methods known in the art can be used to identify binding partner polypeptides, including, (1) organic and inorganic chemical libraries, (2) natural product libraries, and (3) combinatorial libraries comprised of random peptides, oligonucleotides or organic molecules.

The role of downstream intracellular signaling molecules in the signaling cascade of the polypeptide of the invention can be determined. For example, a chimeric protein in which the cytoplasmic domain of the polypeptide of the invention is fused to the extracellular portion of a protein, whose ligand has been identified, is produced in a host cell. The cell is then incubated with the ligand specific for the extracellular portion of the chimeric protein, thereby activating the chimeric receptor. Known downstream proteins involved in intracellular signaling can then be assayed for expected modifications *i.e.* phosphorylation. Other methods known to those in the art can also be used to identify signaling molecules involved in receptor activity.

### 4.10.15 ANTI-INFLAMMATORY ACTIVITY

Compositions of the present invention may also exhibit anti-inflammatory activity. The anti-inflammatory activity may be achieved by providing a stimulus to cells involved in the inflammatory response, by inhibiting or promoting cell-cell interactions (such as, for example, cell adhesion), by inhibiting or promoting chemotaxis of cells involved in the inflammatory process, inhibiting or promoting cell extravasation, or by stimulating or suppressing production of other factors which more directly inhibit or promote an inflammatory response. Compositions with such activities can be used to treat inflammatory conditions including chronic or acute conditions), including without limitation intimation associated with infection (such as septic shock, sepsis or systemic inflammatory response syndrome (SIRS)), ischemia-reperfusion injury, endotoxin lethality, arthritis, complement-mediated hyperacute rejection, nephritis, cytokine or chemokine-induced lung injury, inflammatory bowel disease, Crohn's disease or resulting from over production of cytokines such as TNF or IL-1. Compositions of the invention may also be useful to treat anaphylaxis and hypersensitivity to an antigenic substance or material. Compositions of this invention may be utilized to prevent or treat conditions such as, but not limited to, sepsis, acute pancreatitis, endotoxin shock, cytokine induced shock, rheumatoid arthritis, chronic inflammatory arthritis, pancreatic cell damage from diabetes mellitus type 1, graft versus host disease, inflammatory bowel disease, inflamation associated with pulmonary disease, other autoimmune disease or inflammatory disease, an antiproliferative agent such as for acute or chronic mylegenous leukemia or in the prevention of premature labor secondary to intrauterine infections.

### 4.10.16 LEUKEMIAS

Leukemias and related disorders may be treated or prevented by administration of a therapeutic that promotes or inhibits function of the polynucleotides and/or polypeptides of the invention. Such leukemias and related disorders include but are not limited to acute leukemia, acute lymphocytic leukemia, acute myelocytic leukemia, myeloblastic, promyelocytic, myelomonocytic, monocytic, erythroleukemia, chronic leukemia, chronic myelocytic (granulocytic) leukemia and chronic lymphocytic leukemia (for a review of such disorders, see Fishman et al., 1985, Medicine, 2d Ed., J.B. Lippincott Co., Philadelphia).

### 4.10.17 NERVOUS SYSTEM DISORDERS

Nervous system disorders, involving cell types which can be tested for efficacy of intervention with compounds that modulate the activity of the polynucleotides and/or polypeptides of the invention, and which can be treated upon thus observing an indication of

therapeutic utility, include but are not limited to nervous system injuries, and diseases or disorders which result in either a disconnection of axons, a diminution or degeneration of neurons, or demyelination. Nervous system lesions which may be treated in a patient (including human and non-human mammalian patients) according to the invention include but are not limited to the following lesions of either the central (including spinal cord, brain) or peripheral nervous systems:

(i) traumatic lesions, including lesions caused by physical injury or associated with surgery, for example, lesions which sever a portion of the nervous system, or compression injuries;

(ii) ischemic lesions, in which a lack of oxygen in a portion of the nervous system results in neuronal injury or death, including cerebral infarction or ischemia, or spinal cord infarction or ischemia;

(iii) infectious lesions, in which a portion of the nervous system is destroyed or injured as a result of infection, for example, by an abscess or associated with infection by human immunodeficiency virus, herpes zoster, or herpes simplex virus or with Lyme disease, tuberculosis, syphilis;

(iv) degenerative lesions, in which a portion of the nervous system is destroyed or injured as a result of a degenerative process including but not limited to degeneration associated with Parkinson's disease, Alzheimer's disease, Huntington's chorea, or amyotrophic lateral sclerosis;

(v) lesions associated with nutritional diseases or disorders, in which a portion of the nervous system is destroyed or injured by a nutritional disorder or disorder of metabolism including but not limited to, vitamin B12 deficiency, folic acid deficiency, Wernicke disease, tobacco-alcohol amblyopia, Marchiafava-Bignami disease (primary degeneration of the corpus callosum), and alcoholic cerebellar degeneration;

(vi) neurological lesions associated with systemic diseases including but not limited to diabetes (diabetic neuropathy, Bell's palsy), systemic lupus erythematosus, carcinoma, or sarcoidosis;

(vii) lesions caused by toxic substances including alcohol, lead, or particular neurotoxins; and

(viii) demyelinated lesions in which a portion of the nervous system is destroyed or injured by a demyelinating disease including but not limited to multiple sclerosis, human immunodeficiency virus-associated myelopathy, transverse myelopathy or various etiologies, progressive multifocal leukoencephalopathy, and central pontine myelinolysis.

Therapeutics which are useful according to the invention for treatment of a nervous system disorder may be selected by testing for biological activity in promoting the survival or differentiation of neurons. For example, and not by way of limitation, therapeutics which elicit any of the following effects may be useful according to the invention:

(i) increased survival time of neurons in culture;

(ii) increased sprouting of neurons in culture or in vivo;

(iii) increased production of a neuron-associated molecule in culture or *in vivo*, *e.g.*, choline acetyltransferase or acetylcholinesterase with respect to motor neurons; or

(iv) decreased symptoms of neuron dysfunction in vivo.

Such effects may be measured by any method known in the art. In preferred, non-limiting embodiments, increased survival of neurons may be measured by the method set forth in Arakawa et al. (1990, J. Neurosci. 10:3507-3515); increased sprouting of neurons may be detected by methods set forth in Pestronk et al. (1980, Exp. Neurol. 70:65-82) or Brown et al. (1981, Ann. Rev. Neurosci. 4:17-42); increased production of neuron-associated molecules may be measured by bioassay, enzymatic assay, antibody binding, Northern blot assay, *etc.*, depending on the molecule to be measured; and motor neuron dysfunction may be measured by assessing the physical manifestation of motor neuron disorder, *e.g.*, weakness, motor neuron conduction velocity, or functional disability.

In specific embodiments, motor neuron disorders that may be treated according to the invention include but are not limited to disorders such as infarction, infection, exposure to toxin, trauma, surgical damage, degenerative disease or malignancy that may affect motor neurons as well as other components of the nervous system, as well as disorders that selectively affect neurons such as amyotrophic lateral sclerosis, and including but not limited to progressive spinal muscular atrophy, progressive bulbar palsy, primary lateral sclerosis, infantile and juvenile muscular atrophy, progressive bulbar paralysis of childhood (Fazio-Londe syndrome), poliomyelitis and the post polio syndrome, and Hereditary Motorsensory Neuropathy (Charcot-Marie-Tooth Disease).

### 4.10.18 OTHER ACTIVITIES

A polypeptide of the invention may also exhibit one or more of the following additional activities or effects: inhibiting the growth, infection or function of, or killing, infectious agents, including, without limitation, bacteria, viruses, fungi and other parasites; effecting (suppressing or enhancing) bodily characteristics, including, without limitation, height, weight, hair color, eye color, skin, fat to lean ratio or other tissue pigmentation, or organ or body part size or shape (such as, for example, breast augmentation or diminution, change in bone form or shape);

effecting biorhythms or circadian cycles or rhythms; effecting the fertility of male or female subjects; effecting the metabolism, catabolism, anabolism, processing, utilization, storage or elimination of dietary fat, lipid, protein, carbohydrate, vitamins, minerals, co-factors or other nutritional factors or component(s); effecting behavioral characteristics, including, without limitation, appetite, libido, stress, cognition (including cognitive disorders), depression (including depressive disorders) and violent behaviors; providing analgesic effects or other pain reducing effects; promoting differentiation and growth of embryonic stem cells in lineages other than hematopoietic lineages; hormonal or endocrine activity; in the case of enzymes, correcting deficiencies of the enzyme and treating deficiency-related diseases; treatment of hyperproliferative disorders (such as, for example, psoriasis); immunoglobulin-like activity (such as, for example, the ability to bind antigens or complement); and the ability to act as an antigen in a vaccine composition to raise an immune response against such protein or another material or entity which is cross-reactive with such protein.

### 4.10.19 IDENTIFICATION OF POLYMORPHISMS

The demonstration of polymorphisms makes possible the identification of such polymorphisms in human subjects and the pharmacogenetic use of this information for diagnosis and treatment. Such polymorphisms may be associated with, *e.g.*, differential predisposition or susceptibility to various disease states (such as disorders involving inflammation or immune response) or a differential response to drug administration, and this genetic information can be used to tailor preventive or therapeutic treatment appropriately. For example, the existence of a polymorphism associated with a predisposition to inflammation or autoimmune disease makes possible the diagnosis of this condition in humans by identifying the presence of the polymorphism.

Polymorphisms can be identified in a variety of ways known in the art which all generally involve obtaining a sample from a patient, analyzing DNA from the sample, optionally involving isolation or amplification of the DNA, and identifying the presence of the polymorphism in the DNA. For example, PCR may be used to amplify an appropriate fragment of genomic DNA which may then be sequenced. Alternatively, the DNA may be subjected to allele-specific oligonucleotide hybridization (in which appropriate oligonucleotides are hybridized to the DNA under conditions permitting detection of a single base mismatch) or to a single nucleotide extension assay (in which an oligonucleotide that hybridizes immediately adjacent to the position of the polymorphism is extended with one or more labeled nucleotides). In addition, traditional restriction fragment length polymorphism analysis (using restriction enzymes that provide differential digestion of the genomic DNA depending on the presence or

absence of the polymorphism) may be performed. Arrays with nucleotide sequences of the present invention can be used to detect polymorphisms. The array can comprise modified nucleotide sequences of the present invention in order to detect the nucleotide sequences of the present invention. In the alternative, any one of the nucleotide sequences of the present invention can be placed on the array to detect changes from those sequences.

Alternatively a polymorphism resulting in a change in the amino acid sequence could also be detected by detecting a corresponding change in amino acid sequence of the protein, *e.g.*, by an antibody specific to the variant sequence.

### 4.10.20 ARTHRITIS AND INFLAMMATION

The immunosuppressive effects of the compositions of the invention against rheumatoid arthritis is determined in an experimental animal model system. The experimental model system is adjuvant induced arthritis in rats, and the protocol is described by J. Holoshitz, et at., 1983, Science, 219:56, or by B. Waksman et al., 1963, Int. Arch. Allergy Appl. Immunol., 23:129. Induction of the disease can be caused by a single injection, generally intradermally, of a suspension of killed Mycobacterium tuberculosis in complete Freund's adjuvant (CFA). The route of injection can vary, but rats may be injected at the base of the tail with an adjuvant mixture. The polypeptide is administered in phosphate buffered solution (PBS) at a dose of about 1-5 mg/kg. The control consists of administering PBS only.

The procedure for testing the effects of the test compound would consist of intradermally injecting killed Mycobacterium tuberculosis in CFA followed by immediately administering the test compound and subsequent treatment every other day until day 24. At 14, 15, 18, 20, 22, and 24 days after injection of Mycobacterium CFA, an overall arthritis score may be obtained as described by J. Holoskitz above. An analysis of the data would reveal that the test compound would have a dramatic affect on the swelling of the joints as measured by a decrease of the arthritis score.

## 4.11 THERAPEUTIC METHODS

The compositions (including polypeptide fragments, analogs, variants and antibodies or other binding partners or modulators including antisense polynucleotides) of the invention have numerous applications in a variety of therapeutic methods. Examples of therapeutic applications include, but are not limited to, those exemplified herein.

### 4.11.1 EXAMPLE

One embodiment of the invention is the administration of an effective amount of the polypeptides or other composition of the invention to individuals affected by a disease or disorder that can be modulated by regulating the peptides of the invention. While the mode of administration is not particularly important, parenteral administration is preferred. An exemplary mode of administration is to deliver an intravenous bolus. The dosage of the polypeptides or other composition of the invention will normally be determined by the prescribing physician. It is to be expected that the dosage will vary according to the age, weight, condition and response of the individual patient. Typically, the amount of polypeptide administered per dose will be in the range of about 0.01μg/kg to 100 mg/kg of body weight, with the preferred dose being about 0.1μg/kg to 10 mg/kg of patient body weight. For parenteral administration, polypeptides of the invention will be formulated in an injectable form combined with a pharmaceutically acceptable parenteral vehicle. Such vehicles are well known in the art and examples include water, saline, Ringer's solution, dextrose solution, and solutions consisting of small amounts of the human serum albumin. The vehicle may contain minor amounts of additives that maintain the isotonicity and stability of the polypeptide or other active ingredient. The preparation of such solutions is within the skill of the art.

## 4.12 PHARMACEUTICAL FORMULATIONS AND ROUTES OF ADMINISTRATION

A protein or other composition of the present invention (from whatever source derived, including without limitation from recombinant and non-recombinant sources and including antibodies and other binding partners of the polypeptides of the invention) may be administered to a patient in need, by itself, or in pharmaceutical compositions where it is mixed with suitable carriers or excipient(s) at doses to treat or ameliorate a variety of disorders. Such a composition may optionally contain (in addition to protein or other active ingredient and a carrier) diluents, fillers, salts, buffers, stabilizers, solubilizers, and other materials well known in the art. The term "pharmaceutically acceptable" means a non-toxic material that does not interfere with the effectiveness of the biological activity of the active ingredient(s). The characteristics of the carrier will depend on the route of administration. The pharmaceutical composition of the invention may also contain cytokines, lymphokines, or other hematopoietic factors such as M-CSF, GM-CSF, TNF, IL-1, IL-2, IL-3, IL-4, IL-5, IL-6, IL-7, IL-8, IL-9, IL-10, IL-11, IL-12, IL-13, IL-14, IL-15, IFN, TNF0, TNF1, TNF2, G-CSF, Meg-CSF, thrombopoietin, stem cell factor, and erythropoietin. In further compositions, proteins of the invention may be combined with other agents beneficial to the treatment of the disease or disorder in question. These agents include various growth factors such as epidermal growth factor (EGF), platelet-derived growth

factor (PDGF), transforming growth factors (TGF-α and TGF-β), insulin-like growth factor (IGF), as well as cytokines described herein.

The pharmaceutical composition may further contain other agents which either enhance the activity of the protein or other active ingredient or complement its activity or use in treatment. Such additional factors and/or agents may be included in the pharmaceutical composition to produce a synergistic effect with protein or other active ingredient of the invention, or to minimize side effects. Conversely, protein or other active ingredient of the present invention may be included in formulations of the particular clotting factor, cytokine, lymphokine, other hematopoietic factor, thrombolytic or anti-thrombotic factor, or anti-inflammatory agent to minimize side effects of the clotting factor, cytokine, lymphokine, other hematopoietic factor, thrombolytic or anti-thrombotic factor, or anti-inflammatory agent (such as IL-1Ra, IL-1 Hy1, IL-1 Hy2, anti-TNF, corticosteroids, immunosuppressive agents). A protein of the present invention may be active in multimers (*e.g.*, heterodimers or homodimers) or complexes with itself or other proteins. As a result, pharmaceutical compositions of the invention may comprise a protein of the invention in such multimeric or complexed form.

As an alternative to being included in a pharmaceutical composition of the invention including a first protein, a second protein or a therapeutic agent may be concurrently administered with the first protein (*e.g.*, at the same time, or at differing times provided that therapeutic concentrations of the combination of agents is achieved at the treatment site). Techniques for formulation and administration of the compounds of the instant application may be found in "Remington's Pharmaceutical Sciences," Mack Publishing Co., Easton, PA, latest edition. A therapeutically effective dose further refers to that amount of the compound sufficient to result in amelioration of symptoms, *e.g.*, treatment, healing, prevention or amelioration of the relevant medical condition, or an increase in rate of treatment, healing, prevention or amelioration of such conditions. When applied to an individual active ingredient, administered alone, a therapeutically effective dose refers to that ingredient alone. When applied to a combination, a therapeutically effective dose refers to combined amounts of the active ingredients that result in the therapeutic effect, whether administered in combination, serially or simultaneously.

In practicing the method of treatment or use of the present invention, a therapeutically effective amount of protein or other active ingredient of the present invention is administered to a mammal having a condition to be treated. Protein or other active ingredient of the present invention may be administered in accordance with the method of the invention either alone or in combination with other therapies such as treatments employing cytokines, lymphokines or other hematopoietic factors. When co- administered with one or more cytokines, lymphokines or other

hematopoietic factors, protein or other active ingredient of the present invention may be administered either simultaneously with the cytokine(s), lymphokine(s), other hematopoietic factor(s), thrombolytic or anti-thrombotic factors, or sequentially. If administered sequentially, the attending physician will decide on the appropriate sequence of administering protein or other active ingredient of the present invention in combination with cytokine(s), lymphokine(s), other hematopoietic factor(s), thrombolytic or anti-thrombotic factors.

### 4.12.1 ROUTES OF ADMINISTRATION

Suitable routes of administration may, for example, include oral, rectal, transmucosal, or intestinal administration; parenteral delivery, including intramuscular, subcutaneous, intramedullary injections, as well as intrathecal, direct intraventricular, intravenous, intraperitoneal, intranasal, or intraocular injections. Administration of protein or other active ingredient of the present invention used in the pharmaceutical composition or to practice the method of the present invention can be carried out in a variety of conventional ways, such as oral ingestion, inhalation, topical application or cutaneous, subcutaneous, intraperitoneal, parenteral or intravenous injection. Intravenous administration to the patient is preferred.

Alternately, one may administer the compound in a local rather than systemic manner, for example, via injection of the compound directly into a arthritic joints or in fibrotic tissue, often in a depot or sustained release formulation. In order to prevent the scarring process frequently occurring as complication of glaucoma surgery, the compounds may be administered topically, for example, as eye drops. Furthermore, one may administer the drug in a targeted drug delivery system, for example, in a liposome coated with a specific antibody, targeting, for example, arthritic or fibrotic tissue. The liposomes will be targeted to and taken up selectively by the afflicted tissue.

The polypeptides of the invention are administered by any route that delivers an effective dosage to the desired site of action. The determination of a suitable route of administration and an effective dosage for a particular indication is within the level of skill in the art. Preferably for wound treatment, one administers the therapeutic compound directly to the site. Suitable dosage ranges for the polypeptides of the invention can be extrapolated from these dosages or from similar studies in appropriate animal models. Dosages can then be adjusted as necessary by the clinician to provide maximal therapeutic benefit.

### 4.12.2 COMPOSITIONS/FORMULATIONS

Pharmaceutical compositions for use in accordance with the present invention thus may be formulated in a conventional manner using one or more physiologically acceptable carriers

comprising excipients and auxiliaries which facilitate processing of the active compounds into preparations which can be used pharmaceutically. These pharmaceutical compositions may be manufactured in a manner that is itself known, *e.g.*, by means of conventional mixing, dissolving, granulating, dragee-making, levigating, emulsifying, encapsulating, entrapping or lyophilizing processes. Proper formulation is dependent upon the route of administration chosen. When a therapeutically effective amount of protein or other active ingredient of the present invention is administered orally, protein or other active ingredient of the present invention will be in the form of a tablet, capsule, powder, solution or elixir. When administered in tablet form, the pharmaceutical composition of the invention may additionally contain a solid carrier such as a gelatin or an adjuvant. The tablet, capsule, and powder contain from about 5 to 95% protein or other active ingredient of the present invention, and preferably from about 25 to 90% protein or other active ingredient of the present invention. When administered in liquid form, a liquid carrier such as water, petroleum, oils of animal or plant origin such as peanut oil, mineral oil, soybean oil, or sesame oil, or synthetic oils may be added. The liquid form of the pharmaceutical composition may further contain physiological saline solution, dextrose or other saccharide solution, or glycols such as ethylene glycol, propylene glycol or polyethylene glycol. When administered in liquid form, the pharmaceutical composition contains from about 0.5 to 90% by weight of protein or other active ingredient of the present invention, and preferably from about 1 to 50% protein or other active ingredient of the present invention.

When a therapeutically effective amount of protein or other active ingredient of the present invention is administered by intravenous, cutaneous or subcutaneous injection, protein or other active ingredient of the present invention will be in the form of a pyrogen-free, parenterally acceptable aqueous solution. The preparation of such parenterally acceptable protein or other active ingredient solutions, having due regard to pH, isotonicity, stability, and the like, is within the skill in the art. A preferred pharmaceutical composition for intravenous, cutaneous, or subcutaneous injection should contain, in addition to protein or other active ingredient of the present invention, an isotonic vehicle such as Sodium Chloride Injection, Ringer's Injection, Dextrose Injection, Dextrose and Sodium Chloride Injection, Lactated Ringer's Injection, or other vehicle as known in the art. The pharmaceutical composition of the present invention may also contain stabilizers, preservatives, buffers, antioxidants, or other additives known to those of skill in the art. For injection, the agents of the invention may be formulated in aqueous solutions, preferably in physiologically compatible buffers such as Hanks's solution, Ringer's solution, or physiological saline buffer. For transmucosal administration, penetrants appropriate to the barrier to be permeated are used in the formulation. Such penetrants are generally known in the art.

For oral administration, the compounds can be formulated readily by combining the active compounds with pharmaceutically acceptable carriers well known in the art. Such carriers enable the compounds of the invention to be formulated as tablets, pills, dragees, capsules, liquids, gels, syrups, slurries, suspensions and the like, for oral ingestion by a patient to be treated. Pharmaceutical preparations for oral use can be obtained from a solid excipient, optionally grinding a resulting mixture, and processing the mixture of granules, after adding suitable auxiliaries, if desired, to obtain tablets or dragee cores. Suitable excipients are, in particular, fillers such as sugars, including lactose, sucrose, mannitol, or sorbitol; cellulose preparations such as, for example, maize starch, wheat starch, rice starch, potato starch, gelatin, gum tragacanth, methyl cellulose, hydroxypropylmethyl-cellulose, sodium carboxymethylcellulose, and/or polyvinylpyrrolidone (PVP). If desired, disintegrating agents may be added, such as the cross-linked polyvinyl pyrrolidone, agar, or alginic acid or a salt thereof such as sodium alginate. Dragee cores are provided with suitable coatings. For this purpose, concentrated sugar solutions may be used, which may optionally contain gum arabic, talc, polyvinyl pyrrolidone, carbopol gel, polyethylene glycol, and/or titanium dioxide, lacquer solutions, and suitable organic solvents or solvent mixtures. Dyestuffs or pigments may be added to the tablets or dragee coatings for identification or to characterize different combinations of active compound doses.

Pharmaceutical preparations which can be used orally include push-fit capsules made of gelatin, as well as soft, sealed capsules made of gelatin and a plasticizer, such as glycerol or sorbitol. The push-fit capsules can contain the active ingredients in admixture with filler such as lactose, binders such as starches, and/or lubricants such as talc or magnesium stearate and, optionally, stabilizers. In soft capsules, the active compounds may be dissolved or suspended in suitable liquids, such as fatty oils, liquid paraffin, or liquid polyethylene glycols. In addition, stabilizers may be added. All formulations for oral administration should be in dosages suitable for such administration. For buccal administration, the compositions may take the form of tablets or lozenges formulated in conventional manner.

For administration by inhalation, the compounds for use according to the present invention are conveniently delivered in the form of an aerosol spray presentation from pressurized packs or a nebuliser, with the use of a suitable propellant, *e.g.*, dichlorodifluoromethane, trichlorofluoromethane, dichlorotetrafluoroethane, carbon dioxide or other suitable gas. In the case of a pressurized aerosol the dosage unit may be determined by providing a valve to deliver a metered amount. Capsules and cartridges of, *e.g.*, gelatin for use in an inhaler or insufflator may be formulated containing a powder mix of the compound and a suitable powder base such as lactose or starch. The compounds may be formulated for parenteral

administration by injection, *e.g.*, by bolus injection or continuous infusion. Formulations for injection may be presented in unit dosage form, *e.g.*, in ampules or in multi-dose containers, with an added preservative. The compositions may take such forms as suspensions, solutions or emulsions in oily or aqueous vehicles, and may contain formulatory agents such as suspending, stabilizing and/or dispersing agents.

Pharmaceutical formulations for parenteral administration include aqueous solutions of the active compounds in water-soluble form. Additionally, suspensions of the active compounds may be prepared as appropriate oily injection suspensions. Suitable lipophilic solvents or vehicles include fatty oils such as sesame oil, or synthetic fatty acid esters, such as ethyl oleate or triglycerides, or liposomes. Aqueous injection suspensions may contain substances which increase the viscosity of the suspension, such as sodium carboxymethyl cellulose, sorbitol, or dextran. Optionally, the suspension may also contain suitable stabilizers or agents which increase the solubility of the compounds to allow for the preparation of highly concentrated solutions. Alternatively, the active ingredient may be in powder form for constitution with a suitable vehicle, *e.g.*, sterile pyrogen-free water, before use.

The compounds may also be formulated in rectal compositions such as suppositories or retention enemas, *e.g.*, containing conventional suppository bases such as cocoa butter or other glycerides. In addition to the formulations described previously, the compounds may also be formulated as a depot preparation. Such long acting formulations may be administered by implantation (for example subcutaneously or intramuscularly) or by intramuscular injection. Thus, for example, the compounds may be formulated with suitable polymeric or hydrophobic materials (for example as an emulsion in an acceptable oil) or ion exchange resins, or as sparingly soluble derivatives, for example, as a sparingly soluble salt.

A pharmaceutical carrier for the hydrophobic compounds of the invention is a co-solvent system comprising benzyl alcohol, a nonpolar surfactant, a water-miscible organic polymer, and an aqueous phase. The co-solvent system may be the VPD co-solvent system. VPD is a solution of 3% w/v benzyl alcohol, 8% w/v of the nonpolar surfactant polysorbate 80, and 65% w/v polyethylene glycol 300, made up to volume in absolute ethanol. The VPD co-solvent system (VPD:5W) consists of VPD diluted 1:1 with a 5% dextrose in water solution. This co-solvent system dissolves hydrophobic compounds well, and itself produces low toxicity upon systemic administration. Naturally, the proportions of a co-solvent system may be varied considerably without destroying its solubility and toxicity characteristics. Furthermore, the identity of the co-solvent components may be varied: for example, other low-toxicity nonpolar surfactants may be used instead of polysorbate 80; the fraction size of polyethylene glycol may be varied; other biocompatible polymers may replace polyethylene glycol, *e.g.* polyvinyl pyrrolidone; and other

sugars or polysaccharides may substitute for dextrose. Alternatively, other delivery systems for hydrophobic pharmaceutical compounds may be employed. Liposomes and emulsions are well known examples of delivery vehicles or carriers for hydrophobic drugs. Certain organic solvents such as dimethylsulfoxide also may be employed, although usually at the cost of greater toxicity. Additionally, the compounds may be delivered using a sustained-release system, such as semipermeable matrices of solid hydrophobic polymers containing the therapeutic agent. Various types of sustained-release materials have been established and are well known by those skilled in the art. Sustained-release capsules may, depending on their chemical nature, release the compounds for a few weeks up to over 100 days. Depending on the chemical nature and the biological stability of the therapeutic reagent, additional strategies for protein or other active ingredient stabilization may be employed.

The pharmaceutical compositions also may comprise suitable solid or gel phase carriers or excipients. Examples of such carriers or excipients include but are not limited to calcium carbonate, calcium phosphate, various sugars, starches, cellulose derivatives, gelatin, and polymers such as polyethylene glycols. Many of the active ingredients of the invention may be provided as salts with pharmaceutically compatible counter ions. Such pharmaceutically acceptable base addition salts are those salts which retain the biological effectiveness and properties of the free acids and which are obtained by reaction with inorganic or organic bases such as sodium hydroxide, magnesium hydroxide, ammonia, trialkylamine, dialkylamine, monoalkylamine, dibasic amino acids, sodium acetate, potassium benzoate, triethanol amine and the like.

The pharmaceutical composition of the invention may be in the form of a complex of the protein(s) or other active ingredient(s) of present invention along with protein or peptide antigens. The protein and/or peptide antigen will deliver a stimulatory signal to both B and T lymphocytes. B lymphocytes will respond to antigen through their surface immunoglobulin receptor. T lymphocytes will respond to antigen through the T cell receptor (TCR) following presentation of the antigen by MHC proteins. MHC and structurally related proteins including those encoded by class I and class II MHC genes on host cells will serve to present the peptide antigen(s) to T lymphocytes. The antigen components could also be supplied as purified MHC-peptide complexes alone or with co-stimulatory molecules that can directly signal T cells. Alternatively antibodies able to bind surface immunoglobulin and other molecules on B cells as well as antibodies able to bind the TCR and other molecules on T cells can be combined with the pharmaceutical composition of the invention.

The pharmaceutical composition of the invention may be in the form of a liposome in which protein of the present invention is combined, in addition to other pharmaceutically

acceptable carriers, with amphipathic agents such as lipids which exist in aggregated form as micelles, insoluble monolayers, liquid crystals, or lamellar layers in aqueous solution. Suitable lipids for liposomal formulation include, without limitation, monoglycerides, diglycerides, sulfatides, lysolecithins, phospholipids, saponin, bile acids, and the like. Preparation of such liposomal formulations is within the level of skill in the art, as disclosed, for example, in U.S. Patent Nos. 4,235,871; 4,501,728; 4,837,028; and 4,737,323, all of which are incorporated herein by reference.

The amount of protein or other active ingredient of the present invention in the pharmaceutical composition of the present invention will depend upon the nature and severity of the condition being treated, and on the nature of prior treatments which the patient has undergone. Ultimately, the attending physician will decide the amount of protein or other active ingredient of the present invention with which to treat each individual patient. Initially, the attending physician will administer low doses of protein or other active ingredient of the present invention and observe the patient's response. Larger doses of protein or other active ingredient of the present invention may be administered until the optimal therapeutic effect is obtained for the patient, and at that point the dosage is not increased further. It is contemplated that the various pharmaceutical compositions used to practice the method of the present invention should contain about 0.01 μg to about 100 mg (preferably about 0.1 μg to about 10 mg, more preferably about 0.1 μg to about 1 mg) of protein or other active ingredient of the present invention per kg body weight. For compositions of the present invention which are useful for bone, cartilage, tendon or ligament regeneration, the therapeutic method includes administering the composition topically, systematically, or locally as an implant or device. When administered, the therapeutic composition for use in this invention is, of course, in a pyrogen-free, physiologically acceptable form. Further, the composition may desirably be encapsulated or injected in a viscous form for delivery to the site of bone, cartilage or tissue damage. Topical administration may be suitable for wound healing and tissue repair. Therapeutically useful agents other than a protein or other active ingredient of the invention which may also optionally be included in the composition as described above, may alternatively or additionally, be administered simultaneously or sequentially with the composition in the methods of the invention. Preferably for bone and/or cartilage formation, the composition would include a matrix capable of delivering the protein-containing or other active ingredient-containing composition to the site of bone and/or cartilage damage, providing a structure for the developing bone and cartilage and optimally capable of being resorbed into the body. Such matrices may be formed of materials presently in use for other implanted medical applications.

The choice of matrix material is based on biocompatibility, biodegradability, mechanical properties, cosmetic appearance and interface properties. The particular application of the compositions will define the appropriate formulation. Potential matrices for the compositions may be biodegradable and chemically defined calcium sulfate, tricalcium phosphate, hydroxyapatite, polylactic acid, polyglycolic acid and polyanhydrides. Other potential materials are biodegradable and biologically well-defined, such as bone or dermal collagen. Further matrices are comprised of pure proteins or extracellular matrix components. Other potential matrices are nonbiodegradable and chemically defined, such as sintered hydroxyapatite, bioglass, aluminates, or other ceramics. Matrices may be comprised of combinations of any of the above mentioned types of material, such as polylactic acid and hydroxyapatite or collagen and tricalcium phosphate. The bioceramics may be altered in composition, such as in calcium-aluminate-phosphate and processing to alter pore size, particle size, particle shape, and biodegradability. Presently preferred is a 50:50 (mole weight) copolymer of lactic acid and glycolic acid in the form of porous particles having diameters ranging from 150 to 800 microns. In some applications, it will be useful to utilize a sequestering agent, such as carboxymethyl cellulose or autologous blood clot, to prevent the protein compositions from disassociating from the matrix.

A preferred family of sequestering agents is cellulosic materials such as alkylcelluloses (including hydroxyalkylcelluloses), including methylcellulose, ethylcellulose, hydroxyethylcellulose, hydroxypropylcellulose, hydroxypropyl-methylcellulose, and carboxymethylcellulose, the most preferred being cationic salts of carboxymethylcellulose (CMC). Other preferred sequestering agents include hyaluronic acid, sodium alginate, poly(ethylene glycol), polyoxyethylene oxide, carboxyvinyl polymer and poly(vinyl alcohol). The amount of sequestering agent useful herein is 0.5-20 wt %, preferably 1-10 wt % based on total formulation weight, which represents the amount necessary to prevent desorption of the protein from the polymer matrix and to provide appropriate handling of the composition, yet not so much that the progenitor cells are prevented from infiltrating the matrix, thereby providing the protein the opportunity to assist the osteogenic activity of the progenitor cells. In further compositions, proteins or other active ingredients of the invention may be combined with other agents beneficial to the treatment of the bone and/or cartilage defect, wound, or tissue in question. These agents include various growth factors such as epidermal growth factor (EGF), platelet derived growth factor (PDGF), transforming growth factors (TGF-α and TGF-β), and insulin-like growth factor (IGF).

The therapeutic compositions are also presently valuable for veterinary applications. Particularly domestic animals and thoroughbred horses, in addition to humans, are desired

patients for such treatment with proteins or other active ingredients of the present invention. The dosage regimen of a protein-containing pharmaceutical composition to be used in tissue regeneration will be determined by the attending physician considering various factors which modify the action of the proteins, *e.g.*, amount of tissue weight desired to be formed, the site of damage, the condition of the damaged tissue, the size of a wound, type of damaged tissue (*e.g.*, bone), the patient's age, sex, and diet, the severity of any infection, time of administration and other clinical factors. The dosage may vary with the type of matrix used in the reconstitution and with inclusion of other proteins in the pharmaceutical composition. For example, the addition of other known growth factors, such as IGF I (insulin like growth factor I), to the final composition, may also effect the dosage. Progress can be monitored by periodic assessment of tissue/bone growth and/or repair, for example, X-rays, histomorphometric determinations and tetracycline labeling.

Polynucleotides of the present invention can also be used for gene therapy. Such polynucleotides can be introduced either in vivo or ex vivo into cells for expression in a mammalian subject. Polynucleotides of the invention may also be administered by other known methods for introduction of nucleic acid into a cell or organism (including, without limitation, in the form of viral vectors or naked DNA). Cells may also be cultured ex vivo in the presence of proteins of the present invention in order to proliferate or to produce a desired effect on or activity in such cells. Treated cells can then be introduced in vivo for therapeutic purposes.

### 4.12.3 EFFECTIVE DOSAGE

Pharmaceutical compositions suitable for use in the present invention include compositions wherein the active ingredients are contained in an effective amount to achieve its intended purpose. More specifically, a therapeutically effective amount means an amount effective to prevent development of or to alleviate the existing symptoms of the subject being treated. Determination of the effective amount is well within the capability of those skilled in the art, especially in light of the detailed disclosure provided herein. For any compound used in the method of the invention, the therapeutically effective dose can be estimated initially from appropriate in vitro assays. For example, a dose can be formulated in animal models to achieve a circulating concentration range that can be used to more accurately determine useful doses in humans. For example, a dose can be formulated in animal models to achieve a circulating concentration range that includes the $IC_{50}$ as determined in cell culture (*i.e.*, the concentration of the test compound which achieves a half-maximal inhibition of the protein's biological activity). Such information can be used to more accurately determine useful doses in humans.

A therapeutically effective dose refers to that amount of the compound that results in amelioration of symptoms or a prolongation of survival in a patient. Toxicity and therapeutic efficacy of such compounds can be determined by standard pharmaceutical procedures in cell cultures or experimental animals, *e.g.*, for determining the $LD_{50}$ (the dose lethal to 50% of the population) and the $ED_{50}$ (the dose therapeutically effective in 50% of the population). The dose ratio between toxic and therapeutic effects is the therapeutic index and it can be expressed as the ratio between $LD_{50}$ and $ED_{50}$. Compounds which exhibit high therapeutic indices are preferred. The data obtained from these cell culture assays and animal studies can be used in formulating a range of dosage for use in human. The dosage of such compounds lies preferably within a range of circulating concentrations that include the $ED_{50}$ with little or no toxicity. The dosage may vary within this range depending upon the dosage form employed and the route of administration utilized. The exact formulation, route of administration and dosage can be chosen by the individual physician in view of the patient's condition. See, *e.g.*, Fingl et al., 1975, in "The Pharmacological Basis of Therapeutics", Ch. 1 p.1. Dosage amount and interval may be adjusted individually to provide plasma levels of the active moiety which are sufficient to maintain the desired effects, or minimal effective concentration (MEC). The MEC will vary for each compound but can be estimated from *in vitro* data. Dosages necessary to achieve the MEC will depend on individual characteristics and route of administration. However, HPLC assays or bioassays can be used to determine plasma concentrations.

Dosage intervals can also be determined using MEC value. Compounds should be administered using a regimen which maintains plasma levels above the MEC for 10-90% of the time, preferably between 30-90% and most preferably between 50-90%. In cases of local administration or selective uptake, the effective local concentration of the drug may not be related to plasma concentration.

An exemplary dosage regimen for polypeptides or other compositions of the invention will be in the range of about 0.01 μg/kg to 100 mg/kg of body weight daily, with the preferred dose being about 0.1 μg/kg to 25 mg/kg of patient body weight daily, varying in adults and children. Dosing may be once daily, or equivalent doses may be delivered at longer or shorter intervals.

The amount of composition administered will, of course, be dependent on the subject being treated, on the subject's age and weight, the severity of the affliction, the manner of administration and the judgment of the prescribing physician.

### 4.12.4 PACKAGING

The compositions may, if desired, be presented in a pack or dispenser device which may contain one or more unit dosage forms containing the active ingredient. The pack may, for example, comprise metal or plastic foil, such as a blister pack. The pack or dispenser device may be accompanied by instructions for administration. Compositions comprising a compound of the invention formulated in a compatible pharmaceutical carrier may also be prepared, placed in an appropriate container, and labeled for treatment of an indicated condition.

### 4.13 ANTIBODIES

Also included in the invention are antibodies to proteins, or fragments of proteins of the invention. The term "antibody" as used herein refers to immunoglobulin molecules and immunologically active portions of immunoglobulin (Ig) molecules, *i.e.*, molecules that contain an antigen binding site that specifically binds (immunoreacts with) an antigen. Such antibodies include, but are not limited to, polyclonal, monoclonal, chimeric, single chain, $F_{ab}$, $F_{ab'}$ and $F_{(ab')2}$ fragments, and an $F_{ab}$ expression library. In general, an antibody molecule obtained from humans relates to any of the classes IgG, IgM, IgA, IgE and IgD, which differ from one another by the nature of the heavy chain present in the molecule. Certain classes have subclasses as well, such as $IgG_1$, $IgG_2$, and others. Furthermore, in humans, the light chain may be a kappa chain or a lambda chain. Reference herein to antibodies includes a reference to all such classes, subclasses and types of human antibody species.

An isolated related protein of the invention may be intended to serve as an antigen, or a portion or fragment thereof, and additionally can be used as an immunogen to generate antibodies that immunospecifically bind the antigen, using standard techniques for polyclonal and monoclonal antibody preparation. The full-length protein can be used or, alternatively, the invention provides antigenic peptide fragments of the antigen for use as immunogens. An antigenic peptide fragment comprises at least 6 amino acid residues of the amino acid sequence of the full length protein, (for example the amino acid sequence shown in SEQ ID NO: 1010), and encompasses an epitope thereof such that an antibody raised against the peptide forms a specific immune complex with the full length protein or with any fragment that contains the epitope. Preferably, the antigenic peptide comprises at least 10 amino acid residues, or at least 15 amino acid residues, or at least 20 amino acid residues, or at least 30 amino acid residues. Preferred epitopes encompassed by the antigenic peptide are regions of the protein that are located on its surface; commonly these are hydrophilic regions.

In certain embodiments of the invention, at least one epitope encompassed by the antigenic peptide is a region of -related protein that is located on the surface of the protein, *e.g.*, a hydrophilic region. A hydrophobicity analysis of the human related protein sequence will

indicate which regions of a related protein are particularly hydrophilic and, therefore, are likely to encode surface residues useful for targeting antibody production. As a means for targeting antibody production, hydropathy plots showing regions of hydrophilicity and hydrophobicity may be generated by any method well known in the art, including, for example, the Kyte Doolittle or the Hopp Woods methods, either with or without Fourier transformation. See, *e.g.*, Hopp and Woods, 1981, *Proc. Nat. Acad. Sci. USA* 78: 3824-3828; Kyte and Doolittle 1982, *J. Mol. Biol.* 157: 105-142, each of which is incorporated herein by reference in its entirety. Antibodies that are specific for one or more domains within an antigenic protein, or derivatives, fragments, analogs or homologs thereof, are also provided herein.

A protein of the invention, or a derivative, fragment, analog, homolog or ortholog thereof, may be utilized as an immunogen in the generation of antibodies that immunospecifically bind these protein components.

Various procedures known within the art may be used for the production of polyclonal or monoclonal antibodies directed against a protein of the invention, or against derivatives, fragments, analogs homologs or orthologs thereof (see, for example, Antibodies: A Laboratory Manual, Harlow E, and Lane D, 1988, Cold Spring Harbor Laboratory Press, Cold Spring Harbor, NY, incorporated herein by reference). Some of these antibodies are discussed below.

### 5.13.1 Polyclonal Antibodies

For the production of polyclonal antibodies, various suitable host animals (*e.g.*, rabbit, goat, mouse or other mammal) may be immunized by one or more injections with the native protein, a synthetic variant thereof, or a derivative of the foregoing. An appropriate immunogenic preparation can contain, for example, the naturally occurring immunogenic protein, a chemically synthesized polypeptide representing the immunogenic protein, or a recombinantly expressed immunogenic protein. Furthermore, the protein may be conjugated to a second protein known to be immunogenic in the mammal being immunized. Examples of such immunogenic proteins include but are not limited to keyhole limpet hemocyanin, serum albumin, bovine thyroglobulin, and soybean trypsin inhibitor. The preparation can further include an adjuvant. Various adjuvants used to increase the immunological response include, but are not limited to, Freund's (complete and incomplete), mineral gels (*e.g.*, aluminum hydroxide), surface active substances (*e.g.*, lysolecithin, pluronic polyols, polyanions, peptides, oil emulsions, dinitrophenol, etc.), adjuvants usable in humans such as Bacille Calmette-Guerin and Corynebacterium parvum, or similar immunostimulatory agents. Additional examples of adjuvants which can be employed include MPL-TDM adjuvant (monophosphoryl Lipid A, synthetic trehalose dicorynomycolate).

The polyclonal antibody molecules directed against the immunogenic protein can be isolated from the mammal (*e.g.*, from the blood) and further purified by well known techniques, such as affinity chromatography using protein A or protein G, which provide primarily the IgG fraction of immune serum. Subsequently, or alternatively, the specific antigen which is the target of the immunoglobulin sought, or an epitope thereof, may be immobilized on a column to purify the immune specific antibody by immunoaffinity chromatography. Purification of immunoglobulins is discussed, for example, by D. Wilkinson (The Scientist, published by The Scientist, Inc., Philadelphia PA, Vol. 14, No. 8 (April 17, 2000), pp. 25-28).

### 5.13.2 Monoclonal Antibodies

The term "monoclonal antibody" (MAb) or "monoclonal antibody composition", as used herein, refers to a population of antibody molecules that contain only one molecular species of antibody molecule consisting of a unique light chain gene product and a unique heavy chain gene product. In particular, the complementarity determining regions (CDRs) of the monoclonal antibody are identical in all the molecules of the population. MAbs thus contain an antigen binding site capable of immunoreacting with a particular epitope of the antigen characterized by a unique binding affinity for it.

Monoclonal antibodies can be prepared using hybridoma methods, such as those described by Kohler and Milstein, Nature, 256:495 (1975). In a hybridoma method, a mouse, hamster, or other appropriate host animal, is typically immunized with an immunizing agent to elicit lymphocytes that produce or are capable of producing antibodies that will specifically bind to the immunizing agent. Alternatively, the lymphocytes can be immunized in vitro.

The immunizing agent will typically include the protein antigen, a fragment thereof or a fusion protein thereof. Generally, either peripheral blood lymphocytes are used if cells of human origin are desired, or spleen cells or lymph node cells are used if non-human mammalian sources are desired. The lymphocytes are then fused with an immortalized cell line using a suitable fusing agent, such as polyethylene glycol, to form a hybridoma cell (Goding, Monoclonal Antibodies: Principles and Practice, Academic Press, (1986) pp. 59-103). Immortalized cell lines are usually transformed mammalian cells, particularly myeloma cells of rodent, bovine and human origin. Usually, rat or mouse myeloma cell lines are employed. The hybridoma cells can be cultured in a suitable culture medium that preferably contains one or more substances that inhibit the growth or survival of the unfused, immortalized cells. For example, if the parental cells lack the enzyme hypoxanthine guanine phosphoribosyl transferase (HGPRT or HPRT), the culture medium for the hybridomas typically will include hypoxanthine, aminopterin, and thymidine ("HAT medium"), which substances prevent the growth of HGPRT-deficient cells.

Preferred immortalized cell lines are those that fuse efficiently, support stable high level expression of antibody by the selected antibody-producing cells, and are sensitive to a medium such as HAT medium. More preferred immortalized cell lines are murine myeloma lines, which can be obtained, for instance, from the Salk Institute Cell Distribution Center, San Diego, California and the American Type Culture Collection, Manassas, Virginia. Human myeloma and mouse-human heteromyeloma cell lines also have been described for the production of human monoclonal antibodies (Kozbor, J. Immunol., 133:3001 (1984); Brodeur et al., Monoclonal Antibody Production Techniques and Applications, Marcel Dekker, Inc., New York, (1987) pp. 51-63).

The culture medium in which the hybridoma cells are cultured can then be assayed for the presence of monoclonal antibodies directed against the antigen. Preferably, the binding specificity of monoclonal antibodies produced by the hybridoma cells is determined by immunoprecipitation or by an in vitro binding assay, such as radioimmunoassay (RIA) or enzyme-linked immunoabsorbent assay (ELISA). Such techniques and assays are known in the art. The binding affinity of the monoclonal antibody can, for example, be determined by the Scatchard analysis of Munson and Pollard, Anal. Biochem., 107:220 (1980). Preferably, antibodies having a high degree of specificity and a high binding affinity for the target antigen are isolated.

After the desired hybridoma cells are identified, the clones can be subcloned by limiting dilution procedures and grown by standard methods. Suitable culture media for this purpose include, for example, Dulbecco's Modified Eagle's Medium and RPMI-1640 medium. Alternatively, the hybridoma cells can be grown in vivo as ascites in a mammal.

The monoclonal antibodies secreted by the subclones can be isolated or purified from the culture medium or ascites fluid by conventional immunoglobulin purification procedures such as, for example, protein A-Sepharose, hydroxylapatite chromatography, gel electrophoresis, dialysis, or affinity chromatography.

The monoclonal antibodies can also be made by recombinant DNA methods, such as those described in U.S. Patent No. 4,816,567. DNA encoding the monoclonal antibodies of the invention can be readily isolated and sequenced using conventional procedures (*e.g.*, by using oligonucleotide probes that are capable of binding specifically to genes encoding the heavy and light chains of murine antibodies). The hybridoma cells of the invention serve as a preferred source of such DNA. Once isolated, the DNA can be placed into expression vectors, which are then transfected into host cells such as simian COS cells, Chinese hamster ovary (CHO) cells, or myeloma cells that do not otherwise produce immunoglobulin protein, to obtain the synthesis of monoclonal antibodies in the recombinant host cells. The DNA also can be modified, for

example, by substituting the coding sequence for human heavy and light chain constant domains in place of the homologous murine sequences (U.S. Patent No. 4,816,567; Morrison, Nature 368, 812-13 (1994)) or by covalently joining to the immunoglobulin coding sequence all or part of the coding sequence for a non-immunoglobulin polypeptide. Such a non-immunoglobulin polypeptide can be substituted for the constant domains of an antibody of the invention, or can be substituted for the variable domains of one antigen-combining site of an antibody of the invention to create a chimeric bivalent antibody.

### 5.13.2 Humanized Antibodies

The antibodies directed against the protein antigens of the invention can further comprise humanized antibodies or human antibodies. These antibodies are suitable for administration to humans without engendering an immune response by the human against the administered immunoglobulin. Humanized forms of antibodies are chimeric immunoglobulins, immunoglobulin chains or fragments thereof (such as Fv, Fab, Fab', $F(ab')_2$ or other antigen-binding subsequences of antibodies) that are principally comprised of the sequence of a human immunoglobulin, and contain minimal sequence derived from a non-human immunoglobulin. Humanization can be performed following the method of Winter and co-workers (Jones et al., Nature, 321:522-525 (1986); Riechmann et al., Nature, 332:323-327 (1988); Verhoeyen et al., Science, 239:1534-1536 (1988)), by substituting rodent CDRs or CDR sequences for the corresponding sequences of a human antibody. (See also U.S. Patent No. 5,225,539.) In some instances, Fv framework residues of the human immunoglobulin are replaced by corresponding non-human residues. Humanized antibodies can also comprise residues which are found neither in the recipient antibody nor in the imported CDR or framework sequences. In general, the humanized antibody will comprise substantially all of at least one, and typically two, variable domains, in which all or substantially all of the CDR regions correspond to those of a non-human immunoglobulin and all or substantially all of the framework regions are those of a human immunoglobulin consensus sequence. The humanized antibody optimally also will comprise at least a portion of an immunoglobulin constant region (Fc), typically that of a human immunoglobulin (Jones et al., 1986; Riechmann et al., 1988; and Presta, Curr. Op. Struct. Biol., 2:593-596 (1992)).

### 5.13.3 Human Antibodies

Fully human antibodies relate to antibody molecules in which essentially the entire sequences of both the light chain and the heavy chain, including the CDRs, arise from human genes. Such antibodies are termed "human antibodies", or "fully human antibodies" herein.

Human monoclonal antibodies can be prepared by the trioma technique; the human B-cell hybridoma technique (see Kozbor, et al., 1983 Immunol Today 4: 72) and the EBV hybridoma technique to produce human monoclonal antibodies (see Cole, et al., 1985 In: MONOCLONAL ANTIBODIES AND CANCER THERAPY, Alan R. Liss, Inc., pp. 77-96). Human monoclonal antibodies may be utilized in the practice of the present invention and may be produced by using human hybridomas (see Cote, et al., 1983. Proc Natl Acad Sci USA 80: 2026-2030) or by transforming human B-cells with Epstein Barr Virus in vitro (see Cole, et al., 1985 In: MONOCLONAL ANTIBODIES AND CANCER THERAPY, Alan R. Liss, Inc., pp. 77-96).

In addition, human antibodies can also be produced using additional techniques, including phage display libraries (Hoogenboom and Winter, J. Mol. Biol., 227:381 (1991); Marks et al., J. Mol. Biol., 222:581 (1991)). Similarly, human antibodies can be made by introducing human immunoglobulin loci into transgenic animals, *e.g.*, mice in which the endogenous immunoglobulin genes have been partially or completely inactivated. Upon challenge, human antibody production is observed, which closely resembles that seen in humans in all respects, including gene rearrangement, assembly, and antibody repertoire. This approach is described, for example, in U.S. Patent Nos. 5,545,807; 5,545,806; 5,569,825; 5,625,126; 5,633,425; 5,661,016, and in Marks et al. (Bio/Technology 10, 779-783 (1992)); Lonberg et al. (Nature 368 856-859 (1994)); Morrison ( Nature 368, 812-13 (1994)); Fishwild et al,( Nature Biotechnology 14, 845-51 (1996)); Neuberger (Nature Biotechnology 14, 826 (1996)); and Lonberg and Huszar (Intern. Rev. Immunol. 13 65-93 (1995)).

Human antibodies may additionally be produced using transgenic nonhuman animals which are modified so as to produce fully human antibodies rather than the animal's endogenous antibodies in response to challenge by an antigen. (See PCT publication WO94/02602). The endogenous genes encoding the heavy and light immunoglobulin chains in the nonhuman host have been incapacitated, and active loci encoding human heavy and light chain immunoglobulins are inserted into the host's genome. The human genes are incorporated, for example, using yeast artificial chromosomes containing the requisite human DNA segments. An animal which provides all the desired modifications is then obtained as progeny by crossbreeding intermediate transgenic animals containing fewer than the full complement of the modifications. The preferred embodiment of such a nonhuman animal is a mouse, and is termed the Xenomouse™ as disclosed in PCT publications WO 96/33735 and WO 96/34096. This animal produces B cells which secrete fully human immunoglobulins. The antibodies can be obtained directly from the animal after immunization with an immunogen of interest, as, for example, a preparation of a polyclonal antibody, or alternatively from immortalized B cells derived from the animal, such as hybridomas producing monoclonal antibodies. Additionally, the genes encoding the

immunoglobulins with human variable regions can be recovered and expressed to obtain the antibodies directly, or can be further modified to obtain analogs of antibodies such as, for example, single chain Fv molecules.

An example of a method of producing a nonhuman host, exemplified as a mouse, lacking expression of an endogenous immunoglobulin heavy chain is disclosed in U.S. Patent No. 5,939,598. It can be obtained by a method including deleting the J segment genes from at least one endogenous heavy chain locus in an embryonic stem cell to prevent rearrangement of the locus and to prevent formation of a transcript of a rearranged immunoglobulin heavy chain locus, the deletion being effected by a targeting vector containing a gene encoding a selectable marker; and producing from the embryonic stem cell a transgenic mouse whose somatic and germ cells contain the gene encoding the selectable marker.

A method for producing an antibody of interest, such as a human antibody, is disclosed in U.S. Patent No. 5,916,771. It includes introducing an expression vector that contains a nucleotide sequence encoding a heavy chain into one mammalian host cell in culture, introducing an expression vector containing a nucleotide sequence encoding a light chain into another mammalian host cell, and fusing the two cells to form a hybrid cell. The hybrid cell expresses an antibody containing the heavy chain and the light chain.

In a further improvement on this procedure, a method for identifying a clinically relevant epitope on an immunogen, and a correlative method for selecting an antibody that binds immunospecifically to the relevant epitope with high affinity, are disclosed in PCT publication WO 99/53049.

### 5.13.4 $F_{ab}$ Fragments and Single Chain Antibodies

According to the invention, techniques can be adapted for the production of single-chain antibodies specific to an antigenic protein of the invention (see *e.g.*, U.S. Patent No. 4,946,778). In addition, methods can be adapted for the construction of $F_{ab}$ expression libraries (see *e.g.*, Huse, et al., 1989 Science 246: 1275-1281) to allow rapid and effective identification of monoclonal $F_{ab}$ fragments with the desired specificity for a protein or derivatives, fragments, analogs or homologs thereof. Antibody fragments that contain the idiotypes to a protein antigen may be produced by techniques known in the art including, but not limited to: (i) an $F_{(ab')2}$ fragment produced by pepsin digestion of an antibody molecule; (ii) an $F_{ab}$ fragment generated by reducing the disulfide bridges of an $F_{(ab')2}$ fragment; (iii) an $F_{ab}$ fragment generated by the treatment of the antibody molecule with papain and a reducing agent and (iv) $F_v$ fragments.

### 5.13.5 Bispecific Antibodies

Bispecific antibodies are monoclonal, preferably human or humanized, antibodies that have binding specificities for at least two different antigens. In the present case, one of the binding specificities is for an antigenic protein of the invention. The second binding target is any other antigen, and advantageously is a cell-surface protein or receptor or receptor subunit.

Methods for making bispecific antibodies are known in the art. Traditionally, the recombinant production of bispecific antibodies is based on the co-expression of two immunoglobulin heavy-chain/light-chain pairs, where the two heavy chains have different specificities (Milstein and Cuello, Nature, 305:537-539 (1983)). Because of the random assortment of immunoglobulin heavy and light chains, these hybridomas (quadromas) produce a potential mixture of ten different antibody molecules, of which only one has the correct bispecific structure. The purification of the correct molecule is usually accomplished by affinity chromatography steps. Similar procedures are disclosed in WO 93/08829, published 13 May 1993, and in Traunecker *et al.*, 1991 *EMBO J.*, 10:3655-3659.

Antibody variable domains with the desired binding specificities (antibody-antigen combining sites) can be fused to immunoglobulin constant domain sequences. The fusion preferably is with an immunoglobulin heavy-chain constant domain, comprising at least part of the hinge, CH2, and CH3 regions. It is preferred to have the first heavy-chain constant region (CH1) containing the site necessary for light-chain binding present in at least one of the fusions. DNAs encoding the immunoglobulin heavy-chain fusions and, if desired, the immunoglobulin light chain, are inserted into separate expression vectors, and are co-transfected into a suitable host organism. For further details of generating bispecific antibodies see, for example, Suresh et al., Methods in Enzymology, 121:210 (1986).

According to another approach described in WO 96/27011, the interface between a pair of antibody molecules can be engineered to maximize the percentage of heterodimers which are recovered from recombinant cell culture. The preferred interface comprises at least a part of the CH3 region of an antibody constant domain. In this method, one or more small amino acid side chains from the interface of the first antibody molecule are replaced with larger side chains (*e.g.* tyrosine or tryptophan). Compensatory "cavities" of identical or similar size to the large side chain(s) are created on the interface of the second antibody molecule by replacing large amino acid side chains with smaller ones (*e.g.* alanine or threonine). This provides a mechanism for increasing the yield of the heterodimer over other unwanted end-products such as homodimers.

Bispecific antibodies can be prepared as full length antibodies or antibody fragments (*e.g.* $F(ab')_2$ bispecific antibodies). Techniques for generating bispecific antibodies from antibody fragments have been described in the literature. For example, bispecific antibodies can be prepared using chemical linkage. Brennan et al., Science 229:81 (1985) describe a procedure

wherein intact antibodies are proteolytically cleaved to generate $F(ab')_2$ fragments. These fragments are reduced in the presence of the dithiol complexing agent sodium arsenite to stabilize vicinal dithiols and prevent intermolecular disulfide formation. The Fab' fragments generated are then converted to thionitrobenzoate (TNB) derivatives. One of the Fab'-TNB derivatives is then reconverted to the Fab'-thiol by reduction with mercaptoethylamine and is mixed with an equimolar amount of the other Fab'-TNB derivative to form the bispecific antibody. The bispecific antibodies produced can be used as agents for the selective immobilization of enzymes.

Additionally, Fab' fragments can be directly recovered from E. coli and chemically coupled to form bispecific antibodies. Shalaby et al., J. Exp. Med. 175:217-225 (1992) describe the production of a fully humanized bispecific antibody $F(ab')_2$ molecule. Each Fab' fragment was separately secreted from E. coli and subjected to directed chemical coupling in vitro to form the bispecific antibody. The bispecific antibody thus formed was able to bind to cells overexpressing the ErbB2 receptor and normal human T cells, as well as trigger the lytic activity of human cytotoxic lymphocytes against human breast tumor targets.

Various techniques for making and isolating bispecific antibody fragments directly from recombinant cell culture have also been described. For example, bispecific antibodies have been produced using leucine zippers. Kostelny et al., J. Immunol. 148(5):1547-1553 (1992). The leucine zipper peptides from the Fos and Jun proteins were linked to the Fab' portions of two different antibodies by gene fusion. The antibody homodimers were reduced at the hinge region to form monomers and then re-oxidized to form the antibody heterodimers. This method can also be utilized for the production of antibody homodimers. The "diabody" technology described by Hollinger et al., Proc. Natl. Acad. Sci. USA 90:6444-6448 (1993) has provided an alternative mechanism for making bispecific antibody fragments. The fragments comprise a heavy-chain variable domain ($V_H$) connected to a light-chain variable domain ($V_L$) by a linker which is too short to allow pairing between the two domains on the same chain. Accordingly, the $V_H$ and $V_L$ domains of one fragment are forced to pair with the complementary $V_L$ and $V_H$ domains of another fragment, thereby forming two antigen-binding sites. Another strategy for making bispecific antibody fragments by the use of single-chain Fv (sFv) dimers has also been reported. See, Gruber et al., J. Immunol. 152:5368 (1994).

Antibodies with more than two valencies are contemplated. For example, trispecific antibodies can be prepared. Tutt et al., J. Immunol. 147:60 (1991).

Exemplary bispecific antibodies can bind to two different epitopes, at least one of which originates in the protein antigen of the invention. Alternatively, an anti-antigenic arm of an immunoglobulin molecule can be combined with an arm which binds to a triggering molecule on

a leukocyte such as a T-cell receptor molecule (*e.g.* CD2, CD3, CD28, or B7), or Fc receptors for IgG (FcγR), such as FcγRI (CD64), FcγRII (CD32) and FcγRIII (CD16) so as to focus cellular defense mechanisms to the cell expressing the particular antigen. Bispecific antibodies can also be used to direct cytotoxic agents to cells which express a particular antigen. These antibodies possess an antigen-binding arm and an arm which binds a cytotoxic agent or a radionuclide chelator, such as EOTUBE, DPTA, DOTA, or TETA. Another bispecific antibody of interest binds the protein antigen described herein and further binds tissue factor (TF).

### 5.13.6 Heteroconjugate Antibodies

Heteroconjugate antibodies are also within the scope of the present invention. Heteroconjugate antibodies are composed of two covalently joined antibodies. Such antibodies have, for example, been proposed to target immune system cells to unwanted cells (U.S. Patent No. 4,676,980), and for treatment of HIV infection (WO 91/00360; WO 92/200373; EP 03089). It is contemplated that the antibodies can be prepared in vitro using known methods in synthetic protein chemistry, including those involving crosslinking agents. For example, immunotoxins can be constructed using a disulfide exchange reaction or by forming a thioether bond. Examples of suitable reagents for this purpose include iminothiolate and methyl-4-mercaptobutyrimidate and those disclosed, for example, in U.S. Patent No. 4,676,980.

### 5.13.7 Effector Function Engineering

It can be desirable to modify the antibody of the invention with respect to effector function, so as to enhance, *e.g.*, the effectiveness of the antibody in treating cancer. For example, cysteine residue(s) can be introduced into the Fc region, thereby allowing interchain disulfide bond formation in this region. The homodimeric antibody thus generated can have improved internalization capability and/or increased complement-mediated cell killing and antibody-dependent cellular cytotoxicity (ADCC). See Caron et al., J. Exp Med., 176: 1191-1195 (1992) and Shopes, J. Immunol., 148: 2918-2922 (1992). Homodimeric antibodies with enhanced anti-tumor activity can also be prepared using heterobifunctional cross-linkers as described in Wolff et al. Cancer Research, 53: 2560-2565 (1993). Alternatively, an antibody can be engineered that has dual Fc regions and can thereby have enhanced complement lysis and ADCC capabilities. See Stevenson et al., Anti-Cancer Drug Design, 3: 219-230 (1989).

### 5.13.8 Immunoconjugates

The invention also pertains to immunoconjugates comprising an antibody conjugated to a cytotoxic agent such as a chemotherapeutic agent, toxin (*e.g.*, an enzymatically active toxin of

bacterial, fungal, plant, or animal origin, or fragments thereof), or a radioactive isotope (*i.e.*, a radioconjugate).

Chemotherapeutic agents useful in the generation of such immunoconjugates have been described above. Enzymatically active toxins and fragments thereof that can be used include diphtheria A chain, nonbinding active fragments of diphtheria toxin, exotoxin A chain (from Pseudomonas aeruginosa), ricin A chain, abrin A chain, modeccin A chain, alpha-sarcin, Aleurites fordii proteins, dianthin proteins, Phytolaca americana proteins (PAPI, PAPII, and PAP-S), momordica charantia inhibitor, curcin, crotin, sapaonaria officinalis inhibitor, gelonin, mitogellin, restrictocin, phenomycin, enomycin, and the tricothecenes. A variety of radionuclides are available for the production of radioconjugated antibodies. Examples include $^{212}$Bi, $^{131}$I, $^{131}$In, $^{90}$Y, and $^{186}$Re.

Conjugates of the antibody and cytotoxic agent are made using a variety of bifunctional protein-coupling agents such as N-succinimidyl-3-(2-pyridyldithiol) propionate (SPDP), iminothiolane (IT), bifunctional derivatives of imidoesters (such as dimethyl adipimidate HCL), active esters (such as disuccinimidyl suberate), aldehydes (such as glutareldehyde), bis-azido compounds (such as bis (p-azidobenzoyl) hexanediamine), bis-diazonium derivatives (such as bis-(p-diazoniumbenzoyl)-ethylenediamine), diisocyanates (such as tolyene 2,6-diisocyanate), and bis-active fluorine compounds (such as 1,5-difluoro-2,4-dinitrobenzene). For example, a ricin immunotoxin can be prepared as described in Vitetta et al., Science, 238: 1098 (1987). Carbon-14-labeled 1-isothiocyanatobenzyl-3-methyldiethylene triaminepentaacetic acid (MX-DTPA) is an exemplary chelating agent for conjugation of radionucleotide to the antibody. See WO94/11026.

In another embodiment, the antibody can be conjugated to a "receptor" (such streptavidin) for utilization in tumor pretargeting wherein the antibody-receptor conjugate is administered to the patient, followed by removal of unbound conjugate from the circulation using a clearing agent and then administration of a "ligand" (*e.g.*, avidin) that is in turn conjugated to a cytotoxic agent.

**4.14 COMPUTER READABLE SEQUENCES**

In one application of this embodiment, a nucleotide sequence of the present invention can be recorded on computer readable media. As used herein, "computer readable media" refers to any medium which can be read and accessed directly by a computer. Such media include, but are not limited to: magnetic storage media, such as floppy discs, hard disc storage medium, and magnetic tape; optical storage media such as CD-ROM; electrical storage media such as RAM and ROM; and hybrids of these categories such as magnetic/optical storage media. A skilled

artisan can readily appreciate how any of the presently known computer readable mediums can be used to create a manufacture comprising computer readable medium having recorded thereon a nucleotide sequence of the present invention. As used herein, "recorded" refers to a process for storing information on computer readable medium. A skilled artisan can readily adopt any of the presently known methods for recording information on computer readable medium to generate manufactures comprising the nucleotide sequence information of the present invention.

A variety of data storage structures are available to a skilled artisan for creating a computer readable medium having recorded thereon a nucleotide sequence of the present invention. The choice of the data storage structure will generally be based on the means chosen to access the stored information. In addition, a variety of data processor programs and formats can be used to store the nucleotide sequence information of the present invention on computer readable medium. The sequence information can be represented in a word processing text file, formatted in commercially-available software such as WordPerfect and Microsoft Word, or represented in the form of an ASCII file, stored in a database application, such as DB2, Sybase, Oracle, or the like. A skilled artisan can readily adapt any number of data processor structuring formats (*e.g.* text file or database) in order to obtain computer readable medium having recorded thereon the nucleotide sequence information of the present invention.

By providing any of the nucleotide sequences SEQ ID NO:1-1009 or a representative fragment thereof; or a nucleotide sequence at least 95% identical to any of the nucleotide sequences of SEQ ID NO:1-1009 in computer readable form, a skilled artisan can routinely access the sequence information for a variety of purposes. Computer software is publicly available which allows a skilled artisan to access sequence information provided in a computer readable medium. The examples which follow demonstrate how software which implements the BLAST (Altschul et al., J. Mol. Biol. 215:403-410 (1990)) and BLAZE (Brutlag et al., Comp. Chem. 17:203-207 (1993)) search algorithms on a Sybase system is used to identify open reading frames (ORFs) within a nucleic acid sequence. Such ORFs may be protein encoding fragments and may be useful in producing commercially important proteins such as enzymes used in fermentation reactions and in the production of commercially useful metabolites.

As used herein, "a computer-based system" refers to the hardware means, software means, and data storage means used to analyze the nucleotide sequence information of the present invention. The minimum hardware means of the computer-based systems of the present invention comprises a central processing unit (CPU), input means, output means, and data storage means. A skilled artisan can readily appreciate that any one of the currently available computer-based systems are suitable for use in the present invention. As stated above, the computer-based systems of the present invention comprise a data storage means having stored

therein a nucleotide sequence of the present invention and the necessary hardware means and software means for supporting and implementing a search means. As used herein, "data storage means" refers to memory which can store nucleotide sequence information of the present invention, or a memory access means which can access manufactures having recorded thereon the nucleotide sequence information of the present invention.

As used herein, "search means" refers to one or more programs which are implemented on the computer-based system to compare a target sequence or target structural motif with the sequence information stored within the data storage means. Search means are used to identify fragments or regions of a known sequence which match a particular target sequence or target motif. A variety of known algorithms are disclosed publicly and a variety of commercially available software for conducting search means are and can be used in the computer-based systems of the present invention. Examples of such software includes, but is not limited to, Smith-Waterman, MacPattern (EMBL), BLASTN and BLASTA (NPOLYPEPTIDEIA). A skilled artisan can readily recognize that any one of the available algorithms or implementing software packages for conducting homology searches can be adapted for use in the present computer-based systems. As used herein, a "target sequence" can be any nucleic acid or amino acid sequence of six or more nucleotides or two or more amino acids. A skilled artisan can readily recognize that the longer a target sequence is, the less likely a target sequence will be present as a random occurrence in the database. The most preferred sequence length of a target sequence is from about 10 to 300 amino acids, more preferably from about 30 to 100 nucleotide residues. However, it is well recognized that searches for commercially important fragments, such as sequence fragments involved in gene expression and protein processing, may be of shorter length.

As used herein, "a target structural motif," or "target motif," refers to any rationally selected sequence or combination of sequences in which the sequence(s) are chosen based on a three-dimensional configuration which is formed upon the folding of the target motif. There are a variety of target motifs known in the art. Protein target motifs include, but are not limited to, enzyme active sites and signal sequences. Nucleic acid target motifs include, but are not limited to, promoter sequences, hairpin structures and inducible expression elements (protein binding sequences).

## 4.15 TRIPLE HELIX FORMATION

In addition, the fragments of the present invention, as broadly described, can be used to control gene expression through triple helix formation or antisense DNA or RNA, both of which methods are based on the binding of a polynucleotide sequence to DNA or RNA.

Polynucleotides suitable for use in these methods are preferably 20 to 40 bases in length and are designed to be complementary to a region of the gene involved in transcription (triple helix - see Lee et al., Nucl. Acids Res. 6:3073 (1979); Cooney et al., Science 15241:456 (1988); and Dervan et al., Science 251:1360 (1991)) or to the mRNA itself (antisense - Olmno, J. Neurochem. 56:560 (1991); Oligodeoxynucleotides as Antisense Inhibitors of Gene Expression, CRC Press, Boca Raton, FL (1988)). Triple helix-formation optimally results in a shut-off of RNA transcription from DNA, while antisense RNA hybridization blocks translation of an mRNA molecule into polypeptide. Both techniques have been demonstrated to be effective in model systems. Information contained in the sequences of the present invention is necessary for the design of an antisense or triple helix oligonucleotide.

## 4.16 DIAGNOSTIC ASSAYS AND KITS

The present invention further provides methods to identify the presence or expression of one of the ORFs of the present invention, or homolog thereof, in a test sample, using a nucleic acid probe or antibodies of the present invention, optionally conjugated or otherwise associated with a suitable label.

In general, methods for detecting a polynucleotide of the invention can comprise contacting a sample with a compound that binds to and forms a complex with the polynucleotide for a period sufficient to form the complex, and detecting the complex, so that if a complex is detected, a polynucleotide of the invention is detected in the sample. Such methods can also comprise contacting a sample under stringent hybridization conditions with nucleic acid primers that anneal to a polynucleotide of the invention under such conditions, and amplifying annealed polynucleotides, so that if a polynucleotide is amplified, a polynucleotide of the invention is detected in the sample.

In general, methods for detecting a polypeptide of the invention can comprise contacting a sample with a compound that binds to and forms a complex with the polypeptide for a period sufficient to form the complex, and detecting the complex, so that if a complex is detected, a polypeptide of the invention is detected in the sample.

In detail, such methods comprise incubating a test sample with one or more of the antibodies or one or more of the nucleic acid probes of the present invention and assaying for binding of the nucleic acid probes or antibodies to components within the test sample.

Conditions for incubating a nucleic acid probe or antibody with a test sample vary. Incubation conditions depend on the format employed in the assay, the detection methods employed, and the type and nature of the nucleic acid probe or antibody used in the assay. One skilled in the art will recognize that any one of the commonly available hybridization,

amplification or immunological assay formats can readily be adapted to employ the nucleic acid probes or antibodies of the present invention. Examples of such assays can be found in Chard, T., An Introduction to Radioimmunoassay and Related Techniques, Elsevier Science Publishers, Amsterdam, The Netherlands (1986); Bullock, G.R. et al., Techniques in Immunocytochemistry, Academic Press, Orlando, FL Vol. 1 (1982), Vol. 2 (1983), Vol. 3 (1985); Tijssen, P., Practice and Theory of immunoassays: Laboratory Techniques in Biochemistry and Molecular Biology, Elsevier Science Publishers, Amsterdam, The Netherlands (1985). The test samples of the present invention include cells, protein or membrane extracts of cells, or biological fluids such as sputum, blood, serum, plasma, or urine. The test sample used in the above-described method will vary based on the assay format, nature of the detection method and the tissues, cells or extracts used as the sample to be assayed. Methods for preparing protein extracts or membrane extracts of cells are well known in the art and can be readily be adapted in order to obtain a sample which is compatible with the system utilized.

In another embodiment of the present invention, kits are provided which contain the necessary reagents to carry out the assays of the present invention. Specifically, the invention provides a compartment kit to receive, in close confinement, one or more containers which comprises: (a) a first container comprising one of the probes or antibodies of the present invention; and (b) one or more other containers comprising one or more of the following: wash reagents, reagents capable of detecting presence of a bound probe or antibody.

In detail, a compartment kit includes any kit in which reagents are contained in separate containers. Such containers include small glass containers, plastic containers or strips of plastic or paper. Such containers allows one to efficiently transfer reagents from one compartment to another compartment such that the samples and reagents are not cross-contaminated, and the agents or solutions of each container can be added in a quantitative fashion from one compartment to another. Such containers will include a container which will accept the test sample, a container which contains the antibodies used in the assay, containers which contain wash reagents (such as phosphate buffered saline, Tris-buffers, etc.), and containers which contain the reagents used to detect the bound antibody or probe. Types of detection reagents include labeled nucleic acid probes, labeled secondary antibodies, or in the alternative, if the primary antibody is labeled, the enzymatic, or antibody binding reagents which are capable of reacting with the labeled antibody. One skilled in the art will readily recognize that the disclosed probes and antibodies of the present invention can be readily incorporated into one of the established kit formats which are well known in the art.

## 4.17 MEDICAL IMAGING

The novel polypeptides and binding partners of the invention are useful in medical imaging of sites expressing the molecules of the invention (*e.g.*, where the polypeptide of the invention is involved in the immune response, for imaging sites of inflammation or infection). See, *e.g.*, Kunkel et al., U.S. Pat. NO. 5,413,778. Such methods involve chemical attachment of a labeling or imaging agent, administration of the labeled polypeptide to a subject in a pharmaceutically acceptable carrier, and imaging the labeled polypeptide *in vivo* at the target site.

### 4.18 SCREENING ASSAYS

Using the isolated proteins and polynucleotides of the invention, the present invention further provides methods of obtaining and identifying agents which bind to a polypeptide encoded by an ORF corresponding to any of the nucleotide sequences set forth in SEQ ID NO:1-1009, or bind to a specific domain of the polypeptide encoded by the nucleic acid. In detail, said method comprises the steps of:

(a) contacting an agent with an isolated protein encoded by an ORF of the present invention, or nucleic acid of the invention; and

(b) determining whether the agent binds to said protein or said nucleic acid.

In general, therefore, such methods for identifying compounds that bind to a polynucleotide of the invention can comprise contacting a compound with a polynucleotide of the invention for a time sufficient to form a polynucleotide/compound complex, and detecting the complex, so that if a polynucleotide/compound complex is detected, a compound that binds to a polynucleotide of the invention is identified.

Likewise, in general, therefore, such methods for identifying compounds that bind to a polypeptide of the invention can comprise contacting a compound with a polypeptide of the invention for a time sufficient to form a polypeptide/compound complex, and detecting the complex, so that if a polypeptide/compound complex is detected, a compound that binds to a polynucleotide of the invention is identified.

Methods for identifying compounds that bind to a polypeptide of the invention can also comprise contacting a compound with a polypeptide of the invention in a cell for a time sufficient to form a polypeptide/compound complex, wherein the complex drives expression of a receptor gene sequence in the cell, and detecting the complex by detecting reporter gene sequence expression, so that if a polypeptide/compound complex is detected, a compound that binds a polypeptide of the invention is identified.

Compounds identified via such methods can include compounds which modulate the activity of a polypeptide of the invention (that is, increase or decrease its activity, relative to

activity observed in the absence of the compound). Alternatively, compounds identified via such methods can include compounds which modulate the expression of a polynucleotide of the invention (that is, increase or decrease expression relative to expression levels observed in the absence of the compound). Compounds, such as compounds identified via the methods of the invention, can be tested using standard assays well known to those of skill in the art for their ability to modulate activity/expression.

The agents screened in the above assay can be, but are not limited to, peptides, carbohydrates, vitamin derivatives, or other pharmaceutical agents. The agents can be selected and screened at random or rationally selected or designed using protein modeling techniques.

For random screening, agents such as peptides, carbohydrates, pharmaceutical agents and the like are selected at random and are assayed for their ability to bind to the protein encoded by the ORF of the present invention. Alternatively, agents may be rationally selected or designed. As used herein, an agent is said to be "rationally selected or designed" when the agent is chosen based on the configuration of the particular protein. For example, one skilled in the art can readily adapt currently available procedures to generate peptides, pharmaceutical agents and the like, capable of binding to a specific peptide sequence, in order to generate rationally designed antipeptide peptides, for example see Hurby et al., Application of Synthetic Peptides: Antisense Peptides," In Synthetic Peptides, A User's Guide, W.H. Freeman, NY (1992), pp. 289-307, and Kaspczak et al., Biochemistry 28:9230-8 (1989), or pharmaceutical agents, or the like.

In addition to the foregoing, one class of agents of the present invention, as broadly described, can be used to control gene expression through binding to one of the ORFs or EMFs of the present invention. As described above, such agents can be randomly screened or rationally designed/selected. Targeting the ORF or EMF allows a skilled artisan to design sequence specific or element specific agents, modulating the expression of either a single ORF or multiple ORFs which rely on the same EMF for expression control. One class of DNA binding agents are agents which contain base residues which hybridize or form a triple helix formation by binding to DNA or RNA. Such agents can be based on the classic phosphodiester, ribonucleic acid backbone, or can be a variety of sulfhydryl or polymeric derivatives which have base attachment capacity.

Agents suitable for use in these methods preferably contain 20 to 40 bases and are designed to be complementary to a region of the gene involved in transcription (triple helix - see Lee et al., Nucl. Acids Res. 6:3073 (1979); Cooney et al., Science 241:456 (1988); and Dervan et al., Science 251:1360 (1991)) or to the mRNA itself (antisense - Okano, J. Neurochem. 56:560 (1991); Oligodeoxynucleotides as Antisense Inhibitors of Gene Expression, CRC Press, Boca Raton, FL (1988)). Triple helix-formation optimally results in a shut-off of RNA transcription

from DNA, while antisense RNA hybridization blocks translation of an mRNA molecule into polypeptide. Both techniques have been demonstrated to be effective in model systems. Information contained in the sequences of the present invention is necessary for the design of an antisense or triple helix oligonucleotide and other DNA binding agents.

Agents which bind to a protein encoded by one of the ORFs of the present invention can be used as a diagnostic agent. Agents which bind to a protein encoded by one of the ORFs of the present invention can be formulated using known techniques to generate a pharmaceutical composition.

## 4.19 USE OF NUCLEIC ACIDS AS PROBES

Another aspect of the subject invention is to provide for polypeptide-specific nucleic acid hybridization probes capable of hybridizing with naturally occurring nucleotide sequences. The hybridization probes of the subject invention may be derived from any of the nucleotide sequences SEQ ID NO:1-1009. Because the corresponding gene is only expressed in a limited number of tissues, a hybridization probe derived from of any of the nucleotide sequences SEQ ID NO:1-1009 can be used as an indicator of the presence of RNA of cell type of such a tissue in a sample.

Any suitable hybridization technique can be employed, such as, for example, in situ hybridization. PCR as described in US Patents Nos. 4,683,195 and 4,965,188 provides additional uses for oligonucleotides based upon the nucleotide sequences. Such probes used in PCR may be of recombinant origin, may be chemically synthesized, or a mixture of both. The probe will comprise a discrete nucleotide sequence for the detection of identical sequences or a degenerate pool of possible sequences for identification of closely related genomic sequences.

Other means for producing specific hybridization probes for nucleic acids include the cloning of nucleic acid sequences into vectors for the production of mRNA probes. Such vectors are known in the art and are commercially available and may be used to synthesize RNA probes *in vitro* by means of the addition of the appropriate RNA polymerase as T7 or SP6 RNA polymerase and the appropriate radioactively labeled nucleotides. The nucleotide sequences may be used to construct hybridization probes for mapping their respective genomic sequences. The nucleotide sequence provided herein may be mapped to a chromosome or specific regions of a chromosome using well known genetic and/or chromosomal mapping techniques. These techniques include in situ hybridization, linkage analysis against known chromosomal markers, hybridization screening with libraries or flow-sorted chromosomal preparations specific to known chromosomes, and the like. The technique of fluorescent in situ hybridization of

chromosome spreads has been described, among other places, in Verma et al (1988) Human Chromosomes: A Manual of Basic Techniques, Pergamon Press, New York NY.

Fluorescent *in situ* hybridization of chromosomal preparations and other physical chromosome mapping techniques may be correlated with additional genetic map data. Examples of genetic map data can be found in the 1994 Genome Issue of Science (265:1981f). Correlation between the location of a nucleic acid on a physical chromosomal map and a specific disease (or predisposition to a specific disease) may help delimit the region of DNA associated with that genetic disease. The nucleotide sequences of the subject invention may be used to detect differences in gene sequences between normal, carrier or affected individuals.

### 4.20 PREPARATION OF SUPPORT BOUND OLIGONUCLEOTIDES

Oligonucleotides, *i.e.*, small nucleic acid segments, may be readily prepared by, for example, directly synthesizing the oligonucleotide by chemical means, as is commonly practiced using an automated oligonucleotide synthesizer.

Support bound oligonucleotides may be prepared by any of the methods known to those of skill in the art using any suitable support such as glass, polystyrene or Teflon. One strategy is to precisely spot oligonucleotides synthesized by standard synthesizers. Immobilization can be achieved using passive adsorption (Inouye & Hondo, (1990) J. Clin. Microbiol. 28(6) 1469-72); using UV light (Nagata *et al.,* 1985; Dahlen *et al.,* 1987; Morrissey & Collins, (1989) Mol. Cell Probes 3(2) 189-207) or by covalent binding of base modified DNA (Keller *et al.,* 1988; 1989); all references being specifically incorporated herein.

Another strategy that may be employed is the use of the strong biotin-streptavidin interaction as a linker. For example, Broude *et al.* (1994) Proc. Natl. Acad. Sci. USA 91(8) 3072-6, describe the use of biotinylated probes, although these are duplex probes, that are immobilized on streptavidin-coated magnetic beads. Streptavidin-coated beads may be purchased from Dynal, Oslo. Of course, this same linking chemistry is applicable to coating any surface with streptavidin. Biotinylated probes may be purchased from various sources, such as, *e.g.*, Operon Technologies ~~(Alameda, CA)~~.

Nunc Laboratories (Naperville, IL) is also selling suitable material that could be used. Nunc Laboratories have developed a method by which DNA can be covalently bound to the microwell surface termed Covalink NH. CovaLink NH is a polystyrene surface grafted with secondary amino groups (>NH) that serve as bridge-heads for further covalent coupling. CovaLink Modules may be purchased from Nunc Laboratories. DNA molecules may be bound to CovaLink exclusively at the 5'-end by a phosphoramidate bond, allowing immobilization of more than 1 pmol of DNA (Rasmussen *et al.,* (1991) Anal. Biochem. 198(1) 138-42).

The use of CovaLink NH strips for covalent binding of DNA molecules at the 5'-end has been described (Rasmussen et al., (1991). In this technology, a phosphoramidate bond is employed (Chu et al., (1983) Nucleic Acids Res. 11(8) 6513-29). This is beneficial as immobilization using only a single covalent bond is preferred. The phosphoramidate bond joins the DNA to the CovaLink NH secondary amino groups that are positioned at the end of spacer arms covalently grafted onto the polystyrene surface through a 2 nm long spacer arm. To link an oligonucleotide to CovaLink NH via an phosphoramidate bond, the oligonucleotide terminus must have a 5'-end phosphate group. It is, perhaps, even possible for biotin to be covalently bound to CovaLink and then streptavidin used to bind the probes.

More specifically, the linkage method includes dissolving DNA in water (7.5 ng/ul) and denaturing for 10 min. at 95°C and cooling on ice for 10 min. Ice-cold 0.1 M 1-methylimidazole, pH 7.0 (1-MeIm$_7$), is then added to a final concentration of 10 mM 1-MeIm$_7$. A ss DNA solution is then dispensed into CovaLink NH strips (75 ul/well) standing on ice.

Carbodiimide 0.2 M 1-ethyl-3-(3-dimethylaminopropyl)-carbodiimide (EDC), dissolved in 10 mM 1-MeIm$_7$, is made fresh and 25 ul added per well. The strips are incubated for 5 hours at 50°C. After incubation the strips are washed using, *e.g.*, Nunc-Immuno Wash; first the wells are washed 3 times, then they are soaked with washing solution for 5 min., and finally they are washed 3 times (where in the washing solution is 0.4 N NaOH, 0.25% SDS heated to 50°C).

It is contemplated that a further suitable method for use with the present invention is that described in PCT Patent Application WO 90/03382 (Southern & Maskos), incorporated herein by reference. This method of preparing an oligonucleotide bound to a support involves attaching a nucleoside 3'-reagent through the phosphate group by a covalent phosphodiester link to aliphatic hydroxyl groups carried by the support. The oligonucleotide is then synthesized on the supported nucleoside and protecting groups removed from the synthetic oligonucleotide chain under standard conditions that do not cleave the oligonucleotide from the support. Suitable reagents include nucleoside phosphoramidite and nucleoside hydrogen phosphorate.

An on-chip strategy for the preparation of DNA probe for the preparation of DNA probe arrays may be employed. For example, addressable laser-activated photodeprotection may be employed in the chemical synthesis of oligonucleotides directly on a glass surface, as described by Fodor *et al.* (1991) Science 251(4995) 767-73, incorporated herein by reference. Probes may also be immobilized on nylon supports as described by Van Ness *et al.* (1991) Nucleic Acids Res. 19(12) 3345-50; or linked to Teflon using the method of Duncan & Cavalier (1988) Anal. Biochem. 169(1) 104-8; all references being specifically incorporated herein.

To link an oligonucleotide to a nylon support, as described by Van Ness *et al.* (1991), requires activation of the nylon surface via alkylation and selective activation of the 5'-amine of oligonucleotides with cyanuric chloride.

One particular way to prepare support bound oligonucleotides is to utilize the light-generated synthesis described by Pease *et al.*, (1994) PNAS USA 91(11) 5022-6, incorporated herein by reference). These authors used current photolithographic techniques to generate arrays of immobilized oligonucleotide probes (DNA chips). These methods, in which light is used to direct the synthesis of oligonucleotide probes in high-density, miniaturized arrays, utilize photolabile 5'-protected *N*-acyl-deoxynucleoside phosphoramidites, surface linker chemistry and versatile combinatorial synthesis strategies. A matrix of 256 spatially defined oligonucleotide probes may be generated in this manner.

## 4.21 PREPARATION OF NUCLEIC ACID FRAGMENTS

The nucleic acids may be obtained from any appropriate source, such as cDNAs, genomic DNA, chromosomal DNA, microdissected chromosome bands, cosmid or YAC inserts, and RNA, including mRNA without any amplification steps. For example, Sambrook *et al.* (1989) describes three protocols for the isolation of high molecular weight DNA from mammalian cells (p. 9.14-9.23).

DNA fragments may be prepared as clones in M13, plasmid or lambda vectors and/or prepared directly from genomic DNA or cDNA by PCR or other amplification methods. Samples may be prepared or dispensed in multiwell plates. About 100-1000 ng of DNA samples may be prepared in 2-500 ml of final volume.

The nucleic acids would then be fragmented by any of the methods known to those of skill in the art including, for example, using restriction enzymes as described at 9.24-9.28 of Sambrook *et al.* (1989), shearing by ultrasound and NaOH treatment.

Low pressure shearing is also appropriate, as described by Schriefer *et al.* (1990) Nucleic Acids Res. 18(24) 7455-6, incorporated herein by reference). In this method, DNA samples are passed through a small French pressure cell at a variety of low to intermediate pressures. A lever device allows controlled application of low to intermediate pressures to the cell. The results of these studies indicate that low-pressure shearing is a useful alternative to sonic and enzymatic DNA fragmentation methods.

One particularly suitable way for fragmenting DNA is contemplated to be that using the two base recognition endonuclease, *Cvi*JI, described by Fitzgerald *et al.* (1992) Nucleic Acids Res. 20(14) 3753-62. These authors described an approach for the rapid fragmentation and fractionation

of DNA into particular sizes that they contemplated to be suitable for shotgun cloning and sequencing.

The restriction endonuclease *Cvi*JI normally cleaves the recognition sequence PuGCPy between the G and C to leave blunt ends. Atypical reaction conditions, which alter the specificity of this enzyme (*Cvi*JI**), yield a quasi-random distribution of DNA fragments form the small molecule pUC19 (2688 base pairs). Fitzgerald *et al.* (1992) quantitatively evaluated the randomness of this fragmentation strategy, using a *Cvi*JI** digest of pUC19 that was size fractionated by a rapid gel filtration method and directly ligated, without end repair, to a lac Z minus M13 cloning vector. Sequence analysis of 76 clones showed that *Cvi*JI** restricts pyGCPy and PuGCPu, in addition to PuGCPy sites, and that new sequence data is accumulated at a rate consistent with random fragmentation.

As reported in the literature, advantages of this approach compared to sonication and agarose gel fractionation include: smaller amounts of DNA are required (0.2-0.5 ug instead of 2-5 ug); and fewer steps are involved (no preligation, end repair, chemical extraction, or agarose gel electrophoresis and elution are needed

Irrespective of the manner in which the nucleic acid fragments are obtained or prepared, it is important to denature the DNA to give single stranded pieces available for hybridization. This is achieved by incubating the DNA solution for 2-5 minutes at 80-90°C. The solution is then cooled quickly to 2°C to prevent renaturation of the DNA fragments before they are contacted with the chip. Phosphate groups must also be removed from genomic DNA by methods known in the art.

### 4.22 PREPARATION OF DNA ARRAYS

Arrays may be prepared by spotting DNA samples on a support such as a nylon membrane. Spotting may be performed by using arrays of metal pins (the positions of which correspond to an array of wells in a microtiter plate) to repeated by transfer of about 20 nl of a DNA solution to a nylon membrane. By offset printing, a density of dots higher than the density of the wells is achieved. One to 25 dots may be accommodated in 1 $mm^2$, depending on the type of label used. By avoiding spotting in some preselected number of rows and columns, separate subsets (subarrays) may be formed. Samples in one subarray may be the same genomic segment of DNA (or the same gene) from different individuals, or may be different, overlapped genomic clones. Each of the subarrays may represent replica spotting of the same samples. In one example, a selected gene segment may be amplified from 64 patients. For each patient, the amplified gene segment may be in one 96-well plate (all 96 wells containing the same sample). A plate for each of the 64 patients is prepared. By using a 96-pin device, all samples may be spotted on one 8 x 12 cm membrane.

Subarrays may contain 64 samples, one from each patient. Where the 96 subarrays are identical, the dot span may be 1 $mm^2$ and there may be a 1 mm space between subarrays.

Another approach is to use membranes or plates (available from NUNC, Naperville, Illinois) which may be partitioned by physical spacers *e.g.* a plastic grid molded over the membrane, the grid being similar to the sort of membrane applied to the bottom of multiwell plates, or hydrophobic strips. A fixed physical spacer is not preferred for imaging by exposure to flat phosphor-storage screens or x-ray films.

The present invention is illustrated in the following examples. Upon consideration of the present disclosure, one of skill in the art will appreciate that many other embodiments and variations may be made in the scope of the present invention. Accordingly, it is intended that the broader aspects of the present invention not be limited to the disclosure of the following examples. The present invention is not to be limited in scope by the exemplified embodiments which are intended as illustrations of single aspects of the invention, and compositions and methods which are functionally equivalent are within the scope of the invention. Indeed, numerous modifications and variations in the practice of the invention are expected to occur to those skilled in the art upon consideration of the present preferred embodiments. Consequently, the only limitations which should be placed upon the scope of the invention are those which appear in the appended claims.

All references cited within the body of the instant specification are hereby incorporated by reference in their entirety.

## 5.0 EXAMPLES

### 5.1 EXAMPLE 1

**<u>Novel Nucleic Acid Sequences Obtained From Various Libraries</u>**

A plurality of novel nucleic acids were obtained from cDNA libraries prepared from various human tissues and in some cases isolated from a genomic library derived from human chromosome using standard PCR, SBH sequence signature analysis and Sanger sequencing techniques. The inserts of the library were amplified with PCR using primers specific for the vector sequences which flank the inserts. Clones from cDNA libraries were spotted on nylon membrane filters and screened with oligonucleotide probes (*e.g.*, 7-mers) to obtain signature sequences. The clones were clustered into groups of similar or identical sequences. Representative clones were selected for sequencing.

In some cases, the 5' sequence of the amplified inserts was then deduced using a typical Sanger sequencing protocol. PCR products were purified and subjected to fluorescent dye terminator cycle sequencing. Single pass gel sequencing was done using a 377 Applied Biosystems

(ABI) sequencer to obtain the novel nucleic acid sequences. In some cases RACE (Random Amplification of cDNA Ends) was performed to further extend the sequence in the 5' direction.

### 5.2 EXAMPLE 2

**Novel Contigs**

The novel contigs of the invention were assembled from sequences that were obtained from a cDNA library by methods described in Example 1 above, and in some cases sequences obtained from one or more public databases. Chromatograms were base called and assembled using a software suite from University of Washington, Seattle containing three applications designated PHRED, PHRAP, and CONSED. The sequences for the resulting nucleic acid contigs are designated as SEQ ID NO: 1-1009 and are provided in the attached Sequence Listing. The contigs were assembled using an EST sequence as a seed. Then a recursive algorithm was used to extend the seed EST into an extended assemblage, by pulling additional sequences from different databases (*i.e.*, Hyseq's database containing EST sequences, dbEST version 114, gb pri 114, and UniGene version 101) that belong to this assemblage. The algorithm terminated when there was no additional sequences from the above databases that would extend the assemblage. Inclusion of component sequences into the assemblage was based on a BLASTN hit to the extending assemblage with BLAST score greater than 300 and percent identity greater than 95%.

The nucleotide sequence within the assembled contigs that codes for signal peptide sequences and their cleavage sites was determined from using Neural Network SignalP V1.1 program (from Center for Biological Sequence Analysis, The Technical University of Denmark). The process for identifying prokaryotic and eukaryotic signal peptides and their cleavage sites are also disclosed by Henrik Nielson, Jacob Engelbrecht, Soren Brunak, and Gunnar von Heijne in the publication "Identification of prokaryotic and eukaryotic signal peptides and prediction of their cleavage sites" Protein Engineering, vol. 10, no. 1, pp.1-6 (1997) incorporated herein by reference,. A maximum S score and a mean S score, as described in the Nielson et al. reference, are obtained from each assembled contig. Table 3 sets forth the nucleotide range for each sequence of SEQ ID NO: 1-1009 that encodes a corresponding amino acid sequence containing the signal peptide sequence and its cleavage site: the maximum S score and the mean S score obtained for each sequence.

A signal peptide or leader peptide is usually a segment of about 15 to 30 amino acids at the N terminus of protein that enables the protein to be targeted to a cell membrane or secreted from a cell. Generally, the signal peptide acts as an export lable and is removed as the protein is secreted in its final form.

The nearest neighbor result for the assembled contig was obtained by a BLASTX version 2.01al 19 MP-Washington University search against Genpept release 120 and Geneseq database (October 12, 2000, update 21 (Derwent)), using BLAST algorithm. The nearest neighbor result showed the closest homologue for each assemblage from Genpept (and contains the translated amino acid sequences for which the assemblage encodes). The nearest neighbor results for SEQ ID NO: 1-1009 are shown in Table 2.

Tables 1, 2 and 3 follow. Table 1 shows the various tissue sources of SEQ ID NO: 1-1009. Table 2 shows the nearest neighbor result for the assembled contig. The nearest neighbor result shows the closest homolog with an identifiable function for each assemblage. Table 3 contains the start and stop nucleotides for the translated amino acid sequence for which each assemblage encodes. Table 3 also provides a correlation between the amino acid sequences set forth in the Sequence Listing, the nucleotide sequences set forth in the Sequence Listing and the SEQ ID NO. in USSN 09/491,404.

# TABLE 1

| TISSUE ORIGIN | RNA SOURCE | HYSEQ LIBRARY NAME | SEQ ID NOS: OF NUCLEOTIDE(S) |
|---|---|---|---|
| adult brain | GIBCO | AB3001 | 31 45 61 78 96 122 126 132 163 169 171-172 175-176 181 203 212 220 222 230 251-252 258 263 267 279 336 343 358 396 400-401 422 428-429 431 437 456 464 487 503 513 524 561 580 583 609 619 682 812 946 958 965 980 983 989 999 |
| adult brain | GIBCO | ABD003 | 5 23 26 28-29 31 34-36 61 74 78 87 111-113 116 122-123 129 139 143 148 159 163 167 175-176 178 181 183 186 201-204 206 208-209 212 214 220 222 228 230 234-235 237 246 249-250 252 255 259 262-264 266-267 279-280 286 329 336 351 358 379 396 422 429 431 437 439 444-445 450 452 456 467-468 479 484 503-504 507 513 523-524 526 533 550 553 559 561-562 578 580 583 636 638 640 683 711 759 764 769 772 799 803 824 830 842 865 885 900 902 906 910 922-924 932-933 941 945 951 955 958 965 971 983-984 989 999 1005 |
| adult brain | Clontech | ABR001 | 81 122 148 181 183 204 207 233 237 250 267 301 346 394 396 437 439 457 505 563 618 653 655 721 764 795 885 942 949 |
| adult brain | Clontech | ABR006 | 148 152 222 257 269 583 640 677 878 |
| adult brain | Clontech | ABR008 | 2 10-11 13-14 19-20 23 28-29 34-35 37 39-40 45 49-50 52 60 73-74 78 83 87-91 94 98 101 109 114-117 122-123 143 145 148-150 152 156 162 168 173-178 181 183 187 189 194 204 206-209 212 214-215 220-221 228 231 233-238 246-247 249-253 255-260 262 266 269-270 272 276 278-281 284 294 301 313 316-320 335 337-338 343 363 372 379 388 390-392 396 400-401 403 405-407 414 417 422-423 425 427-428 433 437 441 443-446 452-453 456 464 467 469 473-479 482 484 487-488 491 497-498 500 502 504-505 507 519-520 523-526 533 544-545 553 555-556 563 570-571 574-576 578-580 583 615 618-619 637-638 643-644 653 655-656 661 663 678 680 689-690 695 699 702 705 717-718 720 722 725-726 742 746 752 754-755 759 761 763-765 767 769 772-774 776 784-789 792 795 799 809-810 812 814-815 817 834 840 842 844-846 852 855-856 858-860 870-873 875 877 885-886 888 890-897 903-904 910 928 930-932 939-942 946-947 951-952 955 957 960 964-965 967 971 975-976 978 986-987 989 992 999 1001 |
| adult brain | Clontech | ABR011 | 214 965 |

# TABLE 1

| TISSUE ORIGIN | RNA SOURCE | HYSEQ LIBRARY NAME | SEQ ID NOS: OF NUCLEOTIDE(S) |
|---|---|---|---|
| adult brain | BioChain | ABR012 | 152 498 |
| adult brain | Invitrogen | ABR013 | 142 207 254 396 442 498 |
| adult brain | Invitrogen | ABT004 | 2 23 31 34 78 96 116 129 141 160 176-177 181 183 202 214 231 233 248 256 258-260 262 278 310 336-337 379 416 437 439 443-444 450 452 454 464 467 479 484 500 504 519 526 553 570 590 619 638 640 647 653 655 678 711 759 764 789 795 799 885 887 892 902 905 907 910 915 922 941-942 955 960 989 999 |
| cultured preadipocytes | Strategene | ADP001 | 17 37 39 74 79 111 129 152 160 200 222 248 252 268 274 358 385 450 456 504 526 571 583 619 633 640 740 803 816 829 842 887 939-940 965 973 977 986 |
| adrenal gland | Clontech | ADR002 | 4 6 19 36 39 49 51-53 74 76 118 122-123 147-148 152 156 160 167 171-172 181 183 204 206 212 223-224 228 233-234 246 249-250 254-255 262 274 278-279 284 287 294 317 336 355 358 366 379 392 401-402 412 417 420 431-432 439 464 470 479-480 484 503-504 506 509 519 524 526-527 541 553 555 561 583 614 619 631 638 646 682 738-739 756 760 764 770 800 802-803 816-817 838 847 852 863 881 887 905-906 910 923 926 932 941 950-951 989 999 1002 |
| adult heart | GIBCO | AHR001 | 6 20 26 29 31 34 37 39 41 46 61 74 78 101 114 116-118 122-124 128 145 147-148 152 155 163 175-176 178 181 183 200 204 206 210 212 215 228 230 234-235 237 246 248-252 255-256 262-263 266-268 272 278 280 282-283 286 294 309 313 350-351 358 370 374 379 391-392 394 397 400-401 409 420 423 431-432 434 436 438 441 443 452 455-456 461 467-468 479-480 484 487 498 500 503 505 511 519 533 541 550 552-553 558 561-562 568 575 583 590 597-598 603 619 636-638 644-645 667-668 680 684 711-712 714-715 723 732 750 789 803 805 816 822 828 885 889 900 902 905 908 910 916-917 923-924 932 935 937 939 941 950 952 954 960 965 974 982 984 987 993 1005 |
| adult kidney | GIBCO | AKD001 | 4 13-14 19-20 23 26-31 37 39 47 49 54 61 64 78 81 87 91 98 101 114 118 122-123 127 129-130 141-143 145 148-149 155-158 160 163 168 171-172 175-176 178-181 183 197-198 200 203-206 208 212 215 221-222 228 230 234 237 241 245-246 250-252 254-257 262-263 265-269 278-279 282-284 286 297 301 |

TABLE 1

| TISSUE ORIGIN | RNA SOURCE | HYSEQ LIBRARY NAME | SEQ ID NOS: OF NUCLEOTIDE(S) |
|---|---|---|---|
| | | | 308 333 336 352-353 358 371-372 379 381 386 391 394 396-397 400-401 405 409 417 420 428-429 431 436-437 443 445 450 456 463-466 468 475 479-480 484 487 495 498-499 503-505 507 511 513 517 523 526 529 533 539 541-542 550 552-553 555 561 570-572 575 577-578 583 587 597 604 606 609 619 636 638 640-642 648 680 682 701 706 714 721 732 740 747 771 792 803 805 809 811-812 829 838 842 862 865 885 889 900 902 905-906 908 910-911 918-921 924 926 928-930 937 939 941-942 950-951 953 955 958 960 963 965 967 976 978-979 982-984 1005 |
| adult kidney | Invitrogen | AKT002 | 19 31 78 81 91 98-99 122 142 145 148 152 158 169 176 248 254 256 262 266 279 296-297 301 321 353 372 401 405 416 420 429-430 441 456 464 498 504 507 523 526 533 541 583 592-597 649 701 791 838 862 868 911 926 933 946-947 958 960 971 |
| adult lung | GIBCO | ALG001 | 19 33 48 61 96 98 101 108 111 114 145 148 179 183 194 198 200 205 212 220 228 234 246 248 250-251 254-255 263 268 277 279 289 298 306 337 343 372 379-380 385 401 405-406 408 410 420 431 440 443 445 449 455 484 499 503 507 513 517 571 590 597 617 636 640 714 732 749-750 805 885 900 905 910 918 941 955 958 960 977 980 1001 1005 |
| lymph node | Clontech | ALN001 | 43 48 53 108 123 136 142 147 160 178 181 183 200 205 228 244 246 250 254 268 270 291 379 399 419 431 440 442 479-480 484 519 533 539 553 559 565 583 616-617 619 636 662 701 740 805 833 910 913 928 941 977 |
| young liver | GIBCO | ALV001 | 19 42 45 61 64 84 98 107 109 122-123 129-130 133 142 148 168-169 178 181 183 200 205 207 227-229 232 238 246-248 250 253-255 262-263 265 268 279 317 336 371 377 392 400 410 431 436-437 443 445 448-450 484 487 513 533 545 559 561 570 578 617 632 638 640 648 680 771 803 816 836-838 885 906 926 940 986 |
| adult liver | Invitrogen | ALV002 | 13-14 26 36 54 64 74 76 109 117 122 179 181 183 187 204 215 221 225 229 232 247-248 250 256-257 275 304 307 315 317 321-322 371 377 379 386 416 420 448-449 457 464 475 479 481 483-484 504 507 526 553 557 570 619 627-629 632 |

**TABLE 1**

| TISSUE ORIGIN | RNA SOURCE | HYSEQ LIBRARY NAME | SEQ ID NOS: OF NUCLEOTIDE(S) |
|---|---|---|---|
| | | | 638 640 653 655 675 680 701 752 768 827 848 865 882 885 889 910 951 955 959 963 967 978 989 999-1000 |
| adult ovary | Invitrogen | AOV001 | 4 12 19 23 28-32 34-37 39 45 48 52 54 60-61 64-65 67 76 78 87 96 98-100 108 111-112 114 116-118 122-123 126 129-130 132-134 137 139 142-145 147-149 152 162-163 169-172 176 178 180-183 187 191-192 197-202 204-206 212 214-217 219-222 228 234-235 237 242 246-248 250-252 254-256 262 265-269 274 279-280 282-284 294 308-309 313 317 336-337 346 358 361 364 371 374 379 391-392 394 396-397 400 408 414 418 420 423 425 428-429 431 435-437 440-441 443-447 450 452 455-459 463-464 467-468 479-480 484 487 492 495 499-500 503 505 512-513 517 519 524 533 539 545 553 555 557-559 561 565-566 568 571 575 577-578 581 583 590 597 605 610 613 616-617 619 636 638 640 645-646 649-650 654 662 671 680 682 694 697 701 711 732 735 739-741 750 753 760 764 771 780 785 789 792 803 806 810 812 821 831-832 838 841-842 879 885 887 900 902 905-906 908-912 917 921-922 924 928 936-939 941-942 946 950-952 957-958 960 962-965 979 982 987 989 994 998-999 1005 1008 |
| adult placenta | Clontech | APL001 | 122 148 168 181 194 200 248 262 268 317 436 541 561 803 838 911 971 |
| placenta | Invitrogen | APL002 | 38 61 78-79 142 149 176 187 194 206 215 246 252 278 337 346 379 400 456 464 478-479 484 487 504 519 526 553 571 638 640 732 842 910-911 918 941 958 |
| adult spleen | GIBCO | ASP001 | 23 26 39 43 48 61 63 78 87 98 108 110 123 136 142 157 176 178 181 183 197-198 201-202 205-206 213 220 222 228 234 237 244 250-252 254-255 257 263 294 305 320 336-337 354 358 371-372 376 379 397 400 405 410 414 431 437 440 455-456 484 487 498-499 504 506-507 511-512 519 523 526 529 533 539 550 561 565 572 575 583 586 597 616-617 619 621 636 640 687 701 713 732 740 748 803 812 816 835 910 930 939 946 956 958 |
| testis | GIBCO | ATS001 | 20 23 29 61 64 76 114 123 126 143 145 148-149 175 178 182 200 203 206 209 235 248 252 257 263 268 279-281 283-284 333 358 371 391 396 400 418 423 431 438-439 441 |

TABLE 1

| TISSUE ORIGIN | RNA SOURCE | HYSEQ LIBRARY NAME | SEQ ID NOS: OF NUCLEOTIDE(S) |
|---|---|---|---|
| | | | 445 456 479-480 487 490 505 507-508 516-517 521 524 533 550 559 561-562 582 597 606 638 646 676 680 750 772 803 834 877 908 911 914 937-938 950 989 999 |
| adult bladder | Invitrogen | BLD001 | 23 37 77-78 84 160 176 178 181 215 218 248 252 262 274 299 334 351 401 464 474 484 517 543 619 663 692 729 908 910 918 937 941 951 960 962 |
| bone marrow | Clontech | BMD001 | 19 31 39 43 48 52-53 95-96 98 100 108 111-112 114 117 122-123 136 141-142 144-145 147-149 152 161 163 169 181 183 187 194 201 204-205 208 213 222 228 234 241-242 244-246 248-251 254-255 257 267 272 274 282 286 288-289 292 294 313 317 335 337 339 346-347 358 363 365 374 379 391-392 395-398 406 408 414 418 423 428 436 440-442 444-445 456 475 479 484 495 498-500 504 508 511 516 519 526 533 539 541 553 556 559 561 565 571 573 583 597 612 617 619 638 640 646 649 651 677 681 685 707 709-710 721 734 764 771 803 806 811 838 852 858 869 885 908 910 916 922 930 936-937 941 951 965 982 985 989 991 995 999 1005 1008 |
| bone marrow | Clontech | BMD002 | 31 39 43 48 68 71 91 108 122-123 134 136 142 148-150 152 161 169 178 181 194 196 204-205 208 244 246 254 262-263 265 267 272-273 300 320 343 356 363 372 379 405 408 413-414 430-431 436 440-441 454 479 484 486 512-513 517 519 533 553 559 570 583 590 617-619 634 637 651 674 692 793-794 800 803 818 852 880 904 910 930 936 941 950 |
| bone marrow | Clontech | BMD004 | 142 152 254 274 |
| adult colon | Invitrogen | CLN001 | 26 29 48 61 108-109 129-130 144 176 194 215 221 252 401 436 440 450 498 511 533 583 590 616-617 706 764 905 939 955 |
| adult cervix | BioChain | CVX001 | 6 16 19-20 29 35 37 43 45 64 73 75-76 86 92 96-98 100-101 105 108 111 113 122 143 145 147-149 163-165 167 172 174 178 181-183 187 200-201 206 222 234 237-238 242-243 246 248 250-251 253 261-262 265 268 270 274 279 283-284 294 308 343 345 352 365 379 381 391 400 409 420 423-424 428 436 443-444 463-464 473 479-480 484 487 505 508 510-512 516-517 519 523-524 533 539 553-555 558-559 561-562 575 578 583 591 597 619 643 645-646 650 657 671 680 740 764 771 796 803 811 816 865 889 908 |

TABLE 1

| TISSUE ORIGIN | RNA SOURCE | HYSEQ LIBRARY NAME | SEQ ID NOS: OF NUCLEOTIDE(S) |
|---|---|---|---|
| | | | 910 926-927 933 937 941 960 963 965 967-968 977 982 989 999 1008-1009 |
| diaphragm | BioChain | DIA002 | 26 152 499 680 |
| endothelial cells | Strategene | EDT001 | 13-14 19 23 26 30-32 34 39 67 73-74 76 78 91 101 109 114 116 118 129 145 149 152 156 160-161 167 176 180 183 187 197 201 203-204 206 209 215 222 226 228 230 237 246 248 250-252 256-257 262 266 276 279 282-283 286 309 312-313 343 358 372 391-392 394 396 400-401 405 409 413 420 423 429-431 436 438 443-445 450 455-456 479 484 487 498-499 503 507 509 511 513 523 561-562 571 575 583 619 639 646 653 655 680 711 721 729 739 771-772 775 779 795 803 805 834 838-840 885 889 900 905-906 911 917-918 922 924 930 942 946 955 958 960 977-979 982-984 |
| Genomic clones from the short arm of chromosome 8 | Genomic DNA from Genetic Research | EPM001 | 122 148 436 |
| Genomic clones from the short arm of chromosome 8 | Genomic DNA from Genetic Research | EPM003 | 122 148 379 436 |
| Genomic clones from the short arm of chromosome 8 | Genomic DNA from Genetic Research | EPM004 | 122 148 436 |
| Genomic clones from the short arm of chromosome 8 | Genomic DNA from Genetic Research | EPM005 | 148 |
| esophagus | BioChain | ESO002 | 152 178 583 |
| fetal brain | Clontech | FBR001 | 122 148 181 279 284 484 553 575 619 668 911 |
| fetal brain | Clontech | FBR004 | 122 190 212 379 479 484 541 905 922 924 941 950 |
| fetal brain | Clontech | FBR006 | 2 23 31 36 39 42 44 49 52 78 87 114 117 122-123 145 148 176-177 180-181 187 204 208 210 215 220 235 238-239 241 245-246 251 253 256 259 266 270 278 280 286 314 317 337 372 379 392 396 400-401 405-406 410 414 423 428 439-440 443 445 452 467 473 479 484 487 491 497 500 504 517 519 524 526 544 553 556 561 563 568 570-571 573 577 586 619 647 653 655 664-665 680 739 742 746 754 766 772-776 784 795 798 834 840 842 863 878 885 892-893 898-899 910 930 941-942 946 952 965 971 976 987 993 |
| fetal brain | Invitrogen | FBT002 | 19 31 34-35 44-45 78-79 87 96 101 116 129 176 181 204 206 233 235 |

# TABLE 1

| TISSUE ORIGIN | RNA SOURCE | HYSEQ LIBRARY NAME | SEQ ID NOS: OF NUCLEOTIDE(S) |
|---|---|---|---|
| | | | 256-257 259 262 278 280 317 320 337 380 396-397 401 437 443 446 450 453 464 480 484 498-499 504 526 577 591 619 640 664 680 697 710 764 900 902 905 910 958 |
| fetal heart | Invitrogen | FHR001 | 500 910 |
| fetal kidney | Clontech | FKD001 | 39 47 96 98 122-123 148 156 181 200 207 246 268 274 279 283 300 379 411 445 464 468 479 484 506 542 553 561 583 619 680 686 712 747 910 941 |
| fetal kidney | Clontech | FKD002 | 479 484 583 803 910 941 |
| fetal kidney | Invitrogen | FKD007 | 864 |
| fetal lung | Clontech | FLG001 | 64 96 143-144 168 194 206 234 266 335 337 363 500 507 561 619 968 |
| fetal lung | Invitrogen | FLG003 | 3 13-14 55 61 79 122-123 148 160 181 183 194 200 234 248 250 252 266 268 273 289 294 336 358 428 432 436 484 507 510 513-514 533 541 557-558 582-583 597 671 711 764 777 806 811 817 905 933 978 |
| fetal lung | Clontech | FLG004 | 951 |
| fetal liver-spleen | Columbia University | FLS001 | 13-15 19-21 23-26 28-30 32 34 37 39 45 47-49 56 67 72-74 78 84 87 91 96-98 101 103-104 108 111 114 116 122-123 126 129 131 133 142-145 147-149 151-152 156 160-161 166 168-169 172 176 178-179 181 183-185 192-194 197-202 204-206 208 215 221-222 224 228-229 232 234-235 237 246 248-252 254-257 262 266-268 272 274 278-280 282-287 294 313 315 321 333 336-337 343-344 358 372 377-379 386 391-393 397 400-402 404-405 409-410 418 420-421 429 431 436-437 440-441 443 445 448-450 456-457 464 473 475 478-481 483-484 487-488 498 500 503 505 507 509 513 522-523 528 533-534 541 551 553 558 560-562 564-565 570 575 577-578 583 586 590 597 600 605-607 617 619 632 636 638 640 644 646 672 677-680 705 711 729 732 735-738 740 742 748 760 763-764 771-772 792 802-803 805-806 812 816-817 820-821 824-827 834 838 842-843 848 853 861 865 878 885 887 889 900 902 904-906 908 910-911 917 924 926 928 930 934 936-937 941 944 946 950-951 955 958 960 963 965 974-980 982-983 988-990 999 |
| fetal liver-spleen | Columbia University | FLS002 | 4 8 12 15-16 18-21 23-24 26 32 37 39 47 54 61 64 67 71-72 74 76 79 83-84 87 91 96-98 100-104 109 111-113 122-123 129 133 141 145 147-149 152 161 163 169 171-172 174 178-181 183 185 187-188 192-195 198-202 205 207-209 213 215 221-222 229 232 234-235 237 241 |

TABLE 1

| TISSUE ORIGIN | RNA SOURCE | HYSEQ LIBRARY NAME | SEQ ID NOS: OF NUCLEOTIDE(S) |
|---|---|---|---|
| | | | 244-246 248 250 262 265 267-268 270 274 278-280 283-284 290 294 300 311 313-315 317 331 337 341 346 351-352 358 360-361 371-372 377 382 391-393 397 399-401 404-405 410 414 425 429 431 436 440-441 445-446 448-450 453 456 464 473 475 479-480 487 492 498 500 503-504 507 512 517 519 523 526 540 557 561-563 565 574-575 577-578 583 590 597 605-606 608 611 614 616 619 631-634 636-638 640 646 649-650 662 671-673 676-678 682 684 701-702 704-705 711 716 732 735 748 760 762-764 768 771-772 779 790 802 805 815-816 834 838 842 848 865 878-879 883 887-889 903 905-906 910 916-917 922 924 928 930 939 944 946 950 955-956 958 960 965 975 977 982-983 987-988 993-994 998 1004 |
| fetal liver-spleen | Columbia University | FLS003 | 377 732 889 938 |
| fetal liver | Invitrogen | FLV001 | 23 29 39 84 109 194 208 221 232 247-248 278 301 321 336-337 370-371 379 443 448-449 464 475 479-480 498 500 533 550 578 590 632 636 640 678 680 683 751 763 803 882-883 885 887-889 910 921 942 946 951 963 988 |
| fetal liver | Clontech | FLV004 | 37 122 200 232 268 274 377 583 946 |
| fetal muscle | Invitrogen | FMS001 | 29 37 41 64 66 74 148 164 200 202 208-209 252 257 259 262 265 268 274 279 337 346 379 445 480-481 505 507 553 555 561 571 606 640 676 781 801 838 910 926 928 951 957 960 963 965 |
| fetal muscle | Invitrogen | FMS002 | 200 268 274 |
| fetal skin | Invitrogen | FSK001 | 23 29 31 34 49 78 84 87 96 100 112 116 133 143 148 163 168 172 176-177 181 193 199-202 208 215 222 235 240 246 248 252 256-257 262-268 274 280 282 294 309 314 317 322 346 358 371 373-375 379 414 417 419-420 436-437 441 445 454 456 458 479-480 484 499-500 504 507 513 519-520 526 533 539 541 545-547 550 561 565 570-571 575 577 583 590 598-599 619 644 650 665 697 702 706 739 742 744 784 790 792-793 812 816 861 877 889 906 910 918 922 941 949 951-952 955 962 964-965 968 979 983 987 989 999 |
| fetal skin | Invitrogen | FSK002 | 200 257 265 268 274 513 688 |
| fetal spleen | BioChain | FSP001 | 39 431 523 533 617 |
| umbilical cord | BioChain | FUC001 | 19 28-29 34 39 74 96 99 101 111 114 116 122 143 145 148 163 168 175 178 181 183 197 200 205 212 |

TABLE 1

| TISSUE ORIGIN | RNA SOURCE | HYSEQ LIBRARY NAME | SEQ ID NOS: OF NUCLEOTIDE(S) |
|---|---|---|---|
| | | | 222 228 230 237-238 246 248 252-253 255 257 259 262 265 268-269 272 274 282 325 351 379 396 400-401 413 429 441 443 445 452 456-457 467-468 479 484 487 505 513 517 519 523 533 541 553 555 561 571 575 577 583 590 601-602 605-606 619 636 645 680 693 698 711 757 759 764 803 814 816 821 853 885 889 900 906 908 910 924 926 932 937 941 943 946 951-952 955 958 976 987 989 993-994 999 |
| fetal brain | GIBCO | HFB001 | 13-14 19 26 29 31-32 39 44-45 61 67 74 78 88 100 114 122-123 126 129 148 152 163 167 169 171-172 175-176 180-181 187 201-204 206 209 212 215 220 222 227-228 230 233-235 237 246 249 251 258-259 262-263 266 269 279-280 282 284 286 333 337 340 342 355 358 362 366 379 391 394-397 406 422-423 428-429 431 436-437 443-446 450 452 456 467-468 479-480 484 498 504-505 513 517 523 526-527 533 539 541 558-559 561-562 574 580 583 605 619 635 638 643 680 682 708 711 739-740 742 764 776 803 812 823 865 885 900 902 905 910 917 924 928 932 939 941 945 958 960 964-965 974 978-979 984 |
| macrophage | Invitrogen | HMP001 | 152 201 498 983 |
| infant brain | Columbia University | IB2002 | 2 20 23 26 28-29 31 37 39 44 57 74 78-79 111 118 122-123 126 129 143 145 148 155 168-169 175-176 178 181 185-186 191 200-202 208 212 214-215 220 222 224 228 230-231 235 237 239 248-249 252 255-260 262 266-269 272 280 284 286 289 313 323 326 329 346 358 361 379 396 400 412 422-423 428 437 439 443 445 450 452 457 461 467-468 479-480 484 487 490 498 500 504-505 523 526 533 541-542 547 561-562 571 574-575 580 605 635 637 640 647 653 655 678 680 711 733 746 761 764 766 771 776 795 865 885 887 900-901 905 907 910 917 924 930 932 941-942 951 958 960 962 967 974-975 979 982-983 989 993 999 1003-1004 |
| infant brain | Columbia University | IB2003 | 23 31 53 87 107 123 160 175 185 197 202 207 215 222 237 252 256-258 274 284 289 326 358 396 400 437 445 452 462 464 467 487 500 504 526 575 583 590 605 630 653 655 703 733 757 764 795 865 884-885 900 905 919 924 974-975 982 |
| infant brain | Columbia University | IBM002 | 44 169 248 746 764 958 |
| infant brain | Columbia | IBS001 | 76 119 147 149 181 248 329 361 |

TABLE 1

| TISSUE ORIGIN | RNA SOURCE | HYSEQ LIBRARY NAME | SEQ ID NOS: OF NUCLEOTIDE(S) |
|---|---|---|---|
| | University | | 379 764 910 942 951 |
| lung, fibroblast | Strategene | LFB001 | 13-14 26 78 84 91 98 114 122 148 176 197 204 222 246 251 266 379 387 431 437 441 464 479 484 533 553 571 583 619 645-646 711 739 752 910 926 950 965 978 984 |
| lung tumor | Invitrogen | LGT002 | 13-14 19 31-32 34-39 43 48 64 67 74 76 87 93 95-96 101 111-112 116 122-123 134 138 142 144-145 147-148 151-152 160 172 178-179 181-183 187 191-194 197-198 200-202 205 208 210 218 226 228 234 237 246 248 250-252 254-255 257 260-262 265 268 274 277-279 289 301 320-321 333 336 343 352 355 358 366-368 371 374 379 391-392 397 400-401 406 410 414 423 431 436 440-441 455-456 458 463-464 468 478-480 484 487 498 503-504 511 519 526-527 529 533 541 553 557 561 570-571 575 578 581 583-586 588-589 597 606 616 619 636 638 640 648 650 652 657 680 700 705-706 708 716 721-722 729 732 739 744-745 752 762 764 782 795 803 812 816-817 838 863 874 877 906 910-911 922 926 941 951 955 957-958 962-963 968-969 977-978 982-983 996-997 1007 |
| lymphocytes | ATCC | LPC001 | 13-14 35 66 79 95 106-107 112 122-123 149 152 178 181 201 205 246 251-252 267 293 299 358 379 384 400-401 409 415 418 439 443-444 451 456 458 479 484 487 513 533 568 572 575 583 614 619 686 706 721 730-731 739 747 764 789 905 910 941-942 950 965 978-979 1007 |
| leukocyte | GIBCO | LUC001 | 13-14 19 23 30-32 36 39 45 48-49 60-61 63 67 73-74 78-79 81-82 84 87 91 98-99 107-109 111-112 114 122-123 129 142 144-145 148-150 152 170 176 179 181 183 187-188 194 198 201-208 212-213 215 222 228 235 237 241-242 244-246 249-251 254-257 263 267 278-280 282-284 286 289-290 295 302 308-309 313 317 333 337 343 346 356-358 371 379 391-392 394 397 400-401 404 406-410 412-415 423-424 429 431 436 439-441 443-445 450 456 458 479-480 484 487-488 495 498-500 503 505 511-514 519 523 530-533 539 541 555 559 561 565-566 570 572 577-578 583 590 595 597 617 619 633 635-636 639-640 646 660 670 672 677 680-681 698 703 705 729 732 739-740 743 747 750 763-764 771 782 792-793 803-805 809 819 838 857 866-867 885 888 |

**TABLE 1**

| TISSUE ORIGIN | RNA SOURCE | HYSEQ LIBRARY NAME | SEQ ID NOS: OF NUCLEOTIDE(S) |
|---|---|---|---|
| | | | 900 905 910-911 924 926 928 930 941 948 950-953 955 962-963 965 977-979 984 987 989 999 1008 |
| leukocyte | Clontech | LUC003 | 19 26 68 76 96 122 147 152 198 201 205 208 284 317 354 358 430 436 440 479 511 533 541 553 561 583 589 646 698 732 764 766 838 984 |
| melanoma from cell line ATCC #CRL 1424 | Clontech | MEL004 | 8 23 36 69 91 114 122-123 126 148 151 181 202 204 227 246 256-257 265 313 379 391 400 417 466 478-479 487 496 519 521 523 561 570 583 590 669 728 764 784 838 842 910 941 950 965 970 |
| mammary gland | Invitrogen | MMG001 | 4 19 23 26 29 34-39 43 45 48 55 64 66 74 78 87 96-97 114 116 126 129 136 142 149 151 155-156 160 164 168 173 175-176 178 180-181 183 192 197-200 202 204 207-208 215 222 226-228 230 232 235-238 242 246 248 250 252-257 261-262 268 272 274 278 280 301 303 322 329 335 337 343 363 368-371 374 379 381 391 397 400-401 417 426 429 431 437 439-441 443 445 449-450 455 464 475 478-479 484-485 487-488 498-499 504 507 512 517 519 523 526 532-533 553 557 565 570-571 573 575 577-578 590-591 606 617 619 636 640 646 648 663 677-678 680 691 697 702 708 711 732 744 764 792 803 811-813 817 875-877 885 887-888 900 902 905 908 910-911 918 921-922 934 937 939 941-942 946 951 958 960 965 968 983 989 993 999 1003 1008 |
| induced neuron cells | Strategene | NTD001 | 39 122 148 152 181 212 246 266 313 337 358 379 452 467 479 484 519 553 561 583 621-626 680 872 881 910 924 941 |
| retinoid acid induced neuronal cells | Strategene | NTR001 | 37 148 152 168 541 583 |
| neuronal cells | Strategene | NTU001 | 29 37 147 202 221-222 237 246 262 337 361 391 400 429 439 460 487 504 526 541 583 772 816 924 945 965 |
| pituitary gland | Clontech | PIT004 | 391 396 764 |
| placenta | Clontech | PLA003 | 123 183 544 803 |
| prostate | Clontech | PRT001 | 60-61 76 96 122 145-148 153-154 175 178 183 201 204 226 228 235 237 241 245 248 250-251 256 262 265 280 284 324-325 337 397 400 409 436-437 456 464 478 480 487 489-490 492 508 516-517 524 552 561 583 605 722 740 747 849 889 906 924 926 939 958 974 1005 |
| rectum | Invitrogen | REC001 | 26 29 43 48 70 74 80 108 114 135-136 140 168 178-179 208 226 257 |

TABLE 1

| TISSUE ORIGIN | RNA SOURCE | HYSEQ LIBRARY NAME | SEQ ID NOS: OF NUCLEOTIDE(S) |
|---|---|---|---|
| | | | 262 346 348 371 379 411 413 436-437 475 479 484 499 504 517 526 534 548-549 555 570 577-578 606 636 697 729 764 778 793 885 900 906 908 910 937 941 951 965 989 999 |
| salivary gland | Clontech | SAL001 | 7 38 43 74 87 98 112 122 136 142 148 162 169 181 183-185 207 215 228 235 250 254-255 265 280 349-350 394 437 443 464 508 515-516 519 559 598 614 619 658 666-667 680 724 762-763 771 803 816 842 930 933-934 953 |
| salivary gland | Clontech | SALS03 | 48 108 515 617 900 |
| skin fibroblast | ATCC | SFB001 | 39 |
| skin fibroblast | ATCC | SFB002 | 222 803 |
| skin fibroblast | ATCC | SFB003 | 237 |
| small intestine | Clontech | SIN001 | 16 19 29 39 48 56 65 73 96 108 122 136 148 152 155 160 162 165 168 172 181 191 208 234 244 246 266 282 296 379 394 431 440 443 464 479-480 484 519 571 578 583 617 619 648 662 694 703 752 763 806 838 908 910 926 937 941 966 972 976 |
| skeletal muscle | Clontech | SKM001 | 34 112 116 147 149 152 163 167 373 379 484 515 553 561-562 781 838 910 941 |
| spinal cord | Clontech | SPC001 | 19 22 29 31 55 58 70-71 78 122 134 145 148 150 152 159-160 163 166 171 175-176 183 200-201 203-204 220 222 224 235 237 246 248 250 257 262 266-268 279-280 327-328 330 337 343 346 371 379 389 396 416 429-430 437 443 452-453 456 467 475 479 493-494 498 500 502 541 544 553 561 583 619 635-636 638 640 680 682 696 764 785 900 902 910 941 950 982 994 |
| adult spleen | Clontech | SPLc01 | 254 529 701 |
| stomach | Clontech | STO001 | 48 53 72 74 122 142 152 161 178 181 200-202 204 208 240 251 254 265 268 309 347 397 410 437 512 539 550 583 616 636 657 659 720 722 921 |
| thalamus | Clontech | THA002 | 35 53 78 114 123 156 176 181 228 235 246 252 255-256 265 280 329 331 343 379 437 452 457 467 479 484 496 507 519 553 571 593 619 692 723 754 758 764 853 910 925 941 950 967 981 1003 |
| thymus | Clontech | THM001 | 29 78 112 122 148 151 160-161 169 176 180-181 183 188 198 201 204-206 212 250 254 313 374 379 397 412 429 437 446 453 471-472 484 513 521 529 552-553 561 565 619 636 666 708 739 742 764 771 816 |

# TABLE 1

| TISSUE ORIGIN | RNA SOURCE | HYSEQ LIBRARY NAME | SEQ ID NOS: OF NUCLEOTIDE(S) |
|---|---|---|---|
| | | | 838 910 941-942 944 947 958 969 979 982 989 999 1007 |
| thymus | Clontech | THMc02 | 9 19 32 36 63 67 74 78 80 85-86 122-123 138 142 145 147-148 160-161 169 175-176 181 183-184 187 194 198 202 204 208 211 238 244 246 250 252-254 257 262 265 270-271 283-285 317 333 349 359-360 379 400-401 406 413 418 429 431 433 436 440-441 473 479 484 487 512-513 517-518 523 525 529 533 535-537 541 544 553 556 561 565 567-570 572-573 578 583 615-619 636 644 660-661 681 683 687 698 732 739 763-764 783 785 789 807-808 811 816 842 852 864 868-869 900 904 906 910 924 926 930 938 941 965 968 974 979 992 1006-1007 |
| thyroid gland | Clontech | THR001 | 5 10 13-14 19 23 35 37 39 47 59-61 64 74 79 87 100 110 112 117 122-123 133 141-142 145 148 152 156 160 168 181 187 199-202 204-205 207-208 210 220 224-225 228 234-235 237 246-247 251-252 254-256 262 265 267-268 280-281 284 286 301 308 325 332-333 335 337 343 346 363 371 374 378-379 383 394 396-397 400 420 429 431-432 436 445 452 456 464 467-468 474 479-480 484 487 492 499 507 519 522 533 537 550 553 559 561 569 583 619 638 650 653 655 672 678 680 692 705 719 727 748 764 766-767 769 792 797 816 821 854 906 910-911 921 924 926 928 941 946 951 958 960-961 967 971 974-975 978 984 989 999 |
| trachea | Clontech | TRC001 | 43 48 108 112 142 148 168 204 208 212 221-222 254 265 282 286 317 371 382 425 440 501 553 565 910 |
| uterus | Clontech | UTR001 | 1 37 39 62 145 148 163 183 188 200 257 265 268 346 372 405 408 420 431 520 538 561-562 571 640 680 711 842 850-851 885 910 957 |

# TABLE 2

| SEQ ID NO: OF NUCLEOTIDE | ACCESSION NUMBER | SPECIES | DESCRIPTION | SMITH-WATERMAN SCORE | % IDENTITY |
|---|---|---|---|---|---|
| 1 | AF208846 | Homo sapiens | BM-004 | 172 | 43 |
| 2 | Y53871 | Homo sapiens | A human brain-derived signalling factor polypeptide. | 574 | 99 |
| 3 | AE003620 | Drosophila melanogaster | CG8486 gene product | 112 | 33 |
| 4 | AF193807 | Homo sapiens | Rh type B glycoprotein | 1204 | 96 |
| 5 | Y87156 | Homo sapiens | Human secreted protein sequence SEQ ID NO:195. | 89 | 46 |
| 6 | Y71062 | Homo sapiens | Human membrane transport protein, MTRP-7. | 135 | 30 |
| 7 | AB047936 | Macaca fascicularis | hypothetical protein | 81 | 38 |
| 8 | Y36156 | Homo sapiens | Human secreted protein #28. | 158 | 68 |
| 9 | AB040964 | Homo sapiens | KIAA1531 protein | 495 | 100 |
| 10 | U29725 | Homo sapiens | BMK1 alpha kinase | 114 | 35 |
| 11 | X00822 | Gallus gallus | collagen type III | 54 | 52 |
| 12 | Y27868 | Homo sapiens | Human secreted protein encoded by gene No. 107. | 119 | 43 |
| 13 | W74813 | Homo sapiens | Human secreted protein encoded by gene 85 clone HSDFV29. | 722 | 92 |
| 14 | W74813 | Homo sapiens | Human secreted protein encoded by gene 85 clone HSDFV29. | 722 | 92 |
| 15 | AF119851 | Homo sapiens | PRO1722 | 333 | 70 |
| 16 | AF264750 | Homo sapiens | ALR-like protein | 133 | 100 |
| 17 | X91014 | Mus musculus | alpha 1 type XI collagen | 131 | 72 |
| 18 | AF090930 | Homo sapiens | PRO0478 | 109 | 90 |
| 19 | Y86456 | Homo sapiens | Human gene 46-encoded protein fragment, SEQ ID NO:371. | 618 | 95 |
| 20 | AF084535 | Homo sapiens | laforin | 1809 | 100 |
| 21 | Y27585 | Homo sapiens | Human secreted protein encoded by gene No. 19. | 587 | 98 |
| 22 | Z68748 | Caenorhabditis elegans | Similairity to Yeast hypothetical protein YEH4 (SW:YEH4_YEAST)~cDNA EST yk87c11.3 comes from this gene~cDNA EST yk87c11.5 comes | 214 | 37 |

# TABLE 2

| SEQ ID NO: OF NUCLEOTIDE | ACCESSION NUMBER | SPECIES | DESCRIPTION | SMITH-WATERMAN SCORE | % IDENTITY |
|---|---|---|---|---|---|
| | | | from this gene~cDNA EST yk497d5.3 comes from this gene~cDNA EST yk186a5.5 comes from this gene~cDNA EST yk243b10.5 comes from this gene~cDNA EST yk497d5.5 comes from this gene | | |
| 23 | D86973 | Homo sapiens | similar to Yeast translation activator GCN1 (P1:A48126) | 12053 | 100 |
| 24 | Y09945 | Rattus norvegicus | putative integral membrane transport protein | 458 | 50 |
| 25 | U25739 | Mus musculus | YSPL-1 form 1 | 719 | 77 |
| 26 | AK024427 | Homo sapiens | FLJ00016 protein | 668 | 100 |
| 27 | AP001707 | Homo sapiens | human gene for claudin-8, Accession No. AJ250711 | 603 | 100 |
| 28 | U16030 | Brugia malayi | cuticular collagen Bmcol-2 | 78 | 37 |
| 29 | G02479 | Homo sapiens | Human secreted protein, SEQ ID NO: 6560. | 442 | 100 |
| 30 | Y13375 | Homo sapiens | Amino acid sequence of protein PRO262. | 1806 | 99 |
| 31 | AF077226 | Homo sapiens | copine III | 1757 | 65 |
| 32 | W75198 | Homo sapiens | Human secreted protein encoded by gene 3 clone HCEDO84. | 208 | 100 |
| 33 | AF151978 | Homo sapiens | amino acid transporter B0+ | 3436 | 100 |
| 34 | Y66735 | Homo sapiens | Membrane-bound protein PRO1153. | 1006 | 100 |
| 35 | AC003093 | Homo sapiens | OXYSTEROL-BINDING PROTEIN; 45% similarity to P22059 (PID:g129308) | 764 | 60 |
| 36 | AF286861 | Fasciola hepatica | tegumental antigen-like protein | 79 | 30 |
| 37 | AF201945 | Homo sapiens | HNOEL-iso | 2152 | 100 |
| 38 | AF258465 | Homo sapiens | OTRPC4 | 1668 | 99 |
| 39 | AF173003 | Homo sapiens | apoptosis regulator | 2421 | 100 |
| 40 | Y53023 | Homo sapiens | Human secreted protein clone qf662_3 protein sequence SEQ ID | 128 | 41 |

TABLE 2

| SEQ ID NO: OF NUCLEOTIDE | ACCESSION NUMBER | SPECIES | DESCRIPTION | SMITH-WATERMAN SCORE | % IDENTITY |
|---|---|---|---|---|---|
| | | | NO:52. | | |
| 41 | M25750 | Oryctolagus cuniculus | sarcolumenin precursor | 2307 | 97 |
| 42 | G03797 | Homo sapiens | Human secreted protein, SEQ ID NO: 7878. | 186 | 75 |
| 43 | X57805 | Homo sapiens | immunoglobulin lambda light chain | 1102 | 91 |
| 44 | AE003689 | Drosophila melanogaster | CG4596 gene product | 419 | 44 |
| 45 | Y50934 | Homo sapiens | Human fetal brain cDNA clone vc30_1 derived protein #1. | 644 | 100 |
| 46 | Y19562 | Homo sapiens | Amino acid sequence of a human secreted protein. | 80 | 45 |
| 47 | AF016272 | Homo sapiens | Ksp-cadherin | 4263 | 99 |
| 48 | R13111 | Homo sapiens | 1B1 IgG aberrant light chain with duplicated variable region. | 1000 | 92 |
| 49 | AK001636 | Homo sapiens | unnamed protein product | 1630 | 97 |
| 50 | Y65155 | Homo sapiens | Human 5' EST related polypeptide SEQ ID NO:1316. | 78 | 34 |
| 51 | G00471 | Homo sapiens | Human secreted protein, SEQ ID NO: 4552. | 281 | 91 |
| 52 | AJ272050 | Homo sapiens | transcription initiation factor IA protein | 165 | 68 |
| 53 | Y42388 | Homo sapiens | Amino acid sequence of pt127_1. | 668 | 73 |
| 54 | AF193807 | Homo sapiens | Rh type B glycoprotein | 248 | 97 |
| 55 | AF132611 | Homo sapiens | monocarboxylate transporter MCT3 | 139 | 37 |
| 56 | U43940 | Rattus norvegicus | focal adhesion kinase | 141 | 84 |
| 57 | L17318 | Rattus norvegicus | proline-rich proteoglycan | 124 | 37 |
| 58 | G02832 | Homo sapiens | Human secreted protein, SEQ ID NO: 6913. | 132 | 48 |
| 59 | G00357 | Homo sapiens | Human secreted protein, SEQ ID NO: 4438. | 95 | 64 |
| 60 | Y12723 | Homo sapiens | Human 5' EST secreted protein SEQ ID NO:313. | 91 | 50 |
| 61 | Y19450 | Homo sapiens | Amino acid sequence of a human secreted | 406 | 100 |

# TABLE 2

| SEQ ID NO: OF NUCLEOTIDE | ACCESSION NUMBER | SPECIES | DESCRIPTION | SMITH-WATERMAN SCORE | % IDENTITY |
|---|---|---|---|---|---|
| | | | protein. | | |
| 62 | AF156549 | Mus musculus | putative E1-E2 ATPase | 876 | 65 |
| 63 | AL356276 | Homo sapiens | bA367J7.5 (novel Immunoglobulin domain containing protein) | 655 | 84 |
| 64 | AL133105 | Homo sapiens | hypothetical protein | 1783 | 99 |
| 65 | U32189 | Oryctolagus cuniculus | histidine-rich glycoprotein precursor | 73 | 40 |
| 66 | Y91433 | Homo sapiens | Human secreted protein sequence encoded by gene 33 SEQ ID NO:154. | 758 | 98 |
| 67 | W75198 | Homo sapiens | Human secreted protein encoded by gene 3 clone HCEDO84. | 208 | 100 |
| 68 | AF020651 | Homo sapiens | T cell receptor alpha chain variable region | 742 | 93 |
| 69 | AF118086 | Homo sapiens | PRO1992 | 158 | 61 |
| 70 | X52454 | Drosophila melanogaster | rho | 224 | 36 |
| 71 | W40353 | Homo sapiens | Human unspecified protein from US5702907. | 146 | 67 |
| 72 | Y66690 | Homo sapiens | Membrane-bound protein PRO813. | 971 | 98 |
| 73 | AJ002744 | Homo sapiens | UDP-GalNAc:polypeptide N-acetylgalactosaminyltransferase 7 | 1518 | 98 |
| 74 | AC024792 | Caenorhabditis elegans | contains similarity to TR:P78316 | 423 | 36 |
| 75 | AB016088 | Homo sapiens | RNA binding protein | 109 | 32 |
| 76 | Y94953 | Homo sapiens | Human secreted protein clone fy356_14 protein sequence SEQ ID NO:112. | 2484 | 100 |
| 77 | AF107406 | Homo sapiens | GW128 | 74 | 51 |
| 78 | Y13401 | Homo sapiens | Amino acid sequence of protein PRO339. | 1681 | 96 |
| 79 | Y94290 | Homo sapiens | Human myosin heavy chain homologue. | 1819 | 99 |
| 80 | AF007194 | Homo sapiens | mucin | 4875 | 100 |
| 81 | AF229179 | Homo sapiens | kidney-specific membrane protein NX-17 | 949 | 99 |

# TABLE 2

| SEQ ID NO: OF NUCLEOTIDE | ACCESSION NUMBER | SPECIES | DESCRIPTION | SMITH-WATERMAN SCORE | % IDENTITY |
|---|---|---|---|---|---|
| 82 | AL356173 | Neurospora crassa | hypothetical protein | 83 | 29 |
| 83 | G00437 | Homo sapiens | Human secreted protein, SEQ ID NO: 4518. | 87 | 69 |
| 84 | K03036 | Mus musculus | alpha-1 type I procollagen | 114 | 38 |
| 85 | AF233261 | Homo sapiens | otoraplin | 676 | 100 |
| 86 | AF073519 | Homo sapiens | small EDRK-rich factor 1, long isoform | 100 | 45 |
| 87 | AC021640 | Arabidopsis thaliana | putative phosphatidate phosphohydrolase | 387 | 43 |
| 88 | AB040812 | Homo sapiens | protein kinase PAK5 | 1159 | 100 |
| 89 | AL365409 | Homo sapiens | similar to (NP_034322.1\|) sex-determination protein homolog Fem1a | 694 | 100 |
| 90 | U81035 | Rattus norvegicus | ankyrin binding cell adhesion molecule neurofascin | 189 | 63 |
| 91 | W88684 | Homo sapiens | Secreted protein encoded by gene 151 clone HNHED86. | 134 | 65 |
| 92 | Y66734 | Homo sapiens | Membrane-bound protein PRO1097. | 297 | 70 |
| 93 | AB031051 | Homo sapiens | organic anion transporter OATP-E | 283 | 40 |
| 94 | B08976 | Homo sapiens | Human secreted protein sequence encoded by gene 28 SEQ ID NO:133. | 71 | 27 |
| 95 | U83115 | Homo sapiens | non-lens beta gamma-crystallin like protein | 245 | 97 |
| 96 | AF156551 | Mus musculus | putative E1-E2 ATPase | 3779 | 86 |
| 97 | AF062476 | Mus musculus | retinoic acid-responsive protein; STRA6 | 1091 | 74 |
| 98 | Y87072 | Homo sapiens | Human secreted protein sequence SEQ ID NO:111. | 490 | 100 |
| 99 | AF116652 | Homo sapiens | PRO0813 | 1015 | 99 |
| 100 | AF159567 | Homo sapiens | C2H2 (Kruppel-type) zinc finger protein | 2176 | 100 |
| 101 | D25328 | Homo sapiens | platelet-type phosphofructokinase | 109 | 95 |
| 102 | AB018563 | Homo sapiens | TML1 | 98 | 68 |
| 103 | X83107 | Homo sapiens | bmx | 232 | 85 |

# TABLE 2

| SEQ ID NO: OF NUCLEOTIDE | ACCESSION NUMBER | SPECIES | DESCRIPTION | SMITH-WATERMAN SCORE | % IDENTITY |
|---|---|---|---|---|---|
| 104 | U49973 | Homo sapiens | ORF1; MER37; putative transposase similar to pogo element | 131 | 43 |
| 105 | Y86472 | Homo sapiens | Human gene 52-encoded protein fragment, SEQ ID NO:387. | 150 | 54 |
| 106 | AF020276 | Homo sapiens | spinocerebellar ataxia 7 | 96 | 37 |
| 107 | W57901 | Homo sapiens | Protein of clone CT748_2. | 1499 | 96 |
| 108 | R13111 | Homo sapiens | 1B1 IgG aberrant light chain with duplicated variable region. | 1210 | 84 |
| 109 | W50192 | Homo sapiens | Amino acid sequence of salivary protein CON-1. | 95 | 32 |
| 110 | AB046634 | Macaca fascicularis | hypothetical protein | 282 | 75 |
| 111 | AF242432 | Mus musculus | neuronal apoptosis inhibitory protein 6 | 486 | 29 |
| 112 | AB000280 | Rattus norvegicus | peptide/histidine transporter | 2490 | 88 |
| 113 | AF182443 | Rattus norvegicus | F-box protein FBL2 | 597 | 99 |
| 114 | AJ245874 | Homo sapiens | putative ATG/GTP binding protein | 1242 | 100 |
| 115 | AF179828 | Saimiri sciureus | olfactory receptor | 444 | 66 |
| 116 | Y66735 | Homo sapiens | Membrane-bound protein PRO1153. | 1006 | 100 |
| 117 | Y94344 | Homo sapiens | Human cell surface receptor protein #11. | 892 | 90 |
| 118 | AJ238706 | Drosophila melanogaster | monocarboxylate transporter 1 homologue | 226 | 31 |
| 119 | AF180728 | Drosophila melanogaster | sulfate transporter | 312 | 45 |
| 120 | AE004890 | Pseudomonas aeruginosa | L-lactate permease | 534 | 89 |
| 121 | X91837 | Saccharomyces cerevisiae | cell division cycle protein CDC55 | 435 | 98 |
| 122 | U93565 | Homo sapiens | putative p150 | 1911 | 90 |
| 123 | AJ000332 | Homo sapiens | Glucosidase II | 5043 | 99 |
| 124 | AF204674 | Homo sapiens | muscle disease-related protein | 377 | 72 |
| 125 | S58722 | Homo sapiens | X-linked retinopathy protein {C-terminal, clone | 196 | 68 |

# TABLE 2

| SEQ ID NO: OF NUCLEOTIDE | ACCESSION NUMBER | SPECIES | DESCRIPTION | SMITH-WATERMAN SCORE | % IDENTITY |
|---|---|---|---|---|---|
| | | | XEH.8c} | | |
| 126 | S58722 | Homo sapiens | X-linked retinopathy protein {C-terminal, clone XEH.8c} | 196 | 68 |
| 127 | J03848 | Mesocricetus auratus | metallothionein II | 147 | 51 |
| 128 | G02994 | Homo sapiens | Human secreted protein, SEQ ID NO: 7075. | 93 | 64 |
| 129 | AF116238 | Homo sapiens | pseudouridine synthase 1 | 1927 | 99 |
| 130 | G03411 | Homo sapiens | Human secreted protein, SEQ ID NO: 7492. | 183 | 65 |
| 131 | AF222861 | Sus scrofa | type X collagen | 90 | 34 |
| 132 | G03628 | Homo sapiens | Human secreted protein, SEQ ID NO: 7709. | 60 | 66 |
| 133 | Y10529 | Homo sapiens | olfactory receptor | 766 | 61 |
| 134 | AF164612 | Homo sapiens | Gag protein | 125 | 43 |
| 135 | Y12713 | Mus musculus | Pro-Pol-dUTPase polyprotein | 181 | 47 |
| 136 | X57816 | Homo sapiens | immunoglobulin lambda light chain | 550 | 57 |
| 137 | U07808 | Mus musculus | metallothionein IV | 55 | 37 |
| 138 | AB031227 | Pisum sativum | PsAD1 | 68 | 50 |
| 139 | AB035520 | Oryctolagus cuniculus | parchorin | 1324 | 57 |
| 140 | AB007891 | Homo sapiens | KIAA0431 | 117 | 46 |
| 141 | Y00278 | Homo sapiens | Human secreted protein encoded by gene 21. | 234 | 92 |
| 142 | Y68810 | Homo sapiens | A rat heavy chain region and a human hinge region. | 1124 | 92 |
| 143 | M58526 | Homo sapiens | alpha-5 type IV collagen | 4597 | 97 |
| 144 | AF119851 | Homo sapiens | PRO1722 | 192 | 66 |
| 145 | X84908 | Homo sapiens | phosphorylase kinase | 3798 | 97 |
| 146 | Y76155 | Homo sapiens | Human secreted protein encoded by gene 32. | 81 | 52 |
| 147 | U13766 | Murine leukemia virus | gag-pol polyprotein | 735 | 36 |
| 148 | AF034198 | Homo sapiens | IGSF1 | 7154 | 100 |
| 149 | Y94343 | Homo sapiens | Human cell surface receptor protein #10. | 1331 | 100 |
| 150 | Y87211 | Homo sapiens | Human secreted | 759 | 97 |

# TABLE 2

| SEQ ID NO: OF NUCLEOTIDE | ACCESSION NUMBER | SPECIES | DESCRIPTION | SMITH-WATERMAN SCORE | % IDENTITY |
|---|---|---|---|---|---|
| | | | protein sequence SEQ ID NO:250. | | |
| 151 | AJ252258 | human herpesvirus 2 | glycoprotein G-2 | 115 | 30 |
| 152 | V00662 | Homo sapiens | URF 1 (NADH dehydrogenase subunit) | 1283 | 85 |
| 153 | G02872 | Homo sapiens | Human secreted protein, SEQ ID NO: 6953. | 142 | 61 |
| 154 | A23786 | Beta vulgaris | chitinase 1 | 138 | 41 |
| 155 | Z34465 | Zea mays | extensin-like protein | 97 | 36 |
| 156 | X79389 | Homo sapiens | glutathione transferase T1 | 721 | 66 |
| 157 | M22333 | Homo sapiens | unknown protein | 106 | 46 |
| 158 | AL118502 | Homo sapiens | bA371L19.1 (novel protein) | 2471 | 100 |
| 159 | AJ012582 | Homo sapiens | hyperpolarization-activated cation channel HCN2 | 3076 | 100 |
| 160 | D26351 | Homo sapiens | human type 3 inositol 1,4,5-trisphosphate receptor | 8901 | 99 |
| 161 | AF067656 | Homo sapiens | ZW10 interactor Zwint | 951 | 97 |
| 162 | AE003461 | Drosophila melanogaster | CG11300 gene product | 76 | 29 |
| 163 | Y48518 | Homo sapiens | Human breast tumour-associated protein 63. | 355 | 100 |
| 164 | G00517 | Homo sapiens | Human secreted protein, SEQ ID NO: 4598. | 83 | 34 |
| 165 | G03786 | Homo sapiens | Human secreted protein, SEQ ID NO: 7867. | 251 | 53 |
| 166 | Y00765 | Homo sapiens | Prion protein CJAS. | 63 | 37 |
| 167 | Y21050 | Homo sapiens | Human glial fibrillary acidic protein GFAP mutant fragment 59. | 206 | 71 |
| 168 | X74929 | Homo sapiens | Keratin 8 | 1462 | 95 |
| 169 | U29488 | Caenorhabditis elegans | similar to DNAJ protein | 555 | 29 |
| 170 | L27428 | Homo sapiens | reverse transcriptase | 145 | 45 |
| 171 | W19932 | Homo sapiens | Alzheimer's disease protein encoded by DNA from plasmid pGCS55. | 362 | 100 |
| 172 | AF178983 | Homo sapiens | Ras-associated | 497 | 100 |

# TABLE 2

| SEQ ID NO: OF NUCLEOTIDE | ACCESSION NUMBER | SPECIES | DESCRIPTION | SMITH-WATERMAN SCORE | % IDENTITY |
|---|---|---|---|---|---|
| | | | protein Rap1 | | |
| 173 | U70136 | Homo sapiens | megakaryocyte stimulating factor; MSF | 206 | 28 |
| 174 | G00352 | Homo sapiens | Human secreted protein, SEQ ID NO: 4433. | 109 | 64 |
| 175 | U28143 | Gallus gallus | synemin | 1014 | 39 |
| 176 | Y13401 | Homo sapiens | Amino acid sequence of protein PRO339. | 1978 | 96 |
| 177 | AJ243396 | Homo sapiens | voltage-gated sodium channel beta-3 subunit | 947 | 99 |
| 178 | M77812 | Oryctolagus cuniculus | myosin heavy chain | 4079 | 98 |
| 179 | AF200344 | Homo sapiens | aspartyl protease 3 | 956 | 91 |
| 180 | AF200815 | Homo sapiens | FUSED serine/threonine kinase | 1597 | 99 |
| 181 | G03786 | Homo sapiens | Human secreted protein, SEQ ID NO: 7867. | 147 | 83 |
| 182 | Y00313 | Homo sapiens | Human secreted protein encoded by gene 56. | 56 | 29 |
| 183 | X00699 | Homo sapiens | precursor | 583 | 66 |
| 184 | AF269289 | Homo sapiens | unknown | 81 | 32 |
| 185 | G03797 | Homo sapiens | Human secreted protein, SEQ ID NO: 7878. | 176 | 66 |
| 186 | Y20298 | Homo sapiens | Human apolipoprotein E mutant protein fragment 11. | 110 | 34 |
| 187 | AF161437 | Homo sapiens | HSPC319 | 867 | 99 |
| 188 | Y19684 | Homo sapiens | SEQ ID NO 402 from WO9922243. | 124 | 47 |
| 189 | Y74050 | Homo sapiens | Human prostate tumor EST fragment derived protein #237. | 78 | 42 |
| 190 | Y08986 | Brassica napus | oleosin-like protein | 106 | 36 |
| 191 | AF119851 | Homo sapiens | PRO1722 | 173 | 66 |
| 192 | AF116712 | Homo sapiens | PRO2738 | 166 | 50 |
| 193 | AF186084 | Homo sapiens | epidermal growth factor repeat containing protein | 2022 | 85 |
| 194 | M59819 | Homo sapiens | granulocyte colony-stimulating factor receptor | 4232 | 100 |
| 195 | Y86228 | Homo sapiens | Human secreted protein HFXJX44, | 250 | 100 |

TABLE 2

| SEQ ID NO: OF NUCLEOTIDE | ACCESSION NUMBER | SPECIES | DESCRIPTION | SMITH-WATERMAN SCORE | % IDENTITY |
|---|---|---|---|---|---|
| | | | SEQ ID NO:143. | | |
| 196 | Y45382 | Homo sapiens | Human secreted protein fragment encoded from gene 28. | 181 | 63 |
| 197 | X94991 | Homo sapiens | zyxin | 566 | 41 |
| 198 | M17236 | Homo sapiens | MHC HLA-DQ alpha precursor | 896 | 84 |
| 199 | AC004659 | Homo sapiens | BC62940_2 | 805 | 53 |
| 200 | X14420 | Homo sapiens | prepro-alpha-1 type 3 collagen | 5521 | 99 |
| 201 | AF180473 | Homo sapiens | Not2p | 1628 | 98 |
| 202 | X85237 | Homo sapiens | human splicing factor | 1145 | 100 |
| 203 | AL390114 | Leishmania major | extremely cysteine/valine rich protein | 309 | 58 |
| 204 | D42138 | Homo sapiens | PIG-B | 1479 | 98 |
| 205 | Y00062 | Homo sapiens | precursor polypeptide (AA -23 to 1120) | 3334 | 98 |
| 206 | W93946 | Homo sapiens | Human regulatory molecule HRM-2 protein. | 1011 | 100 |
| 207 | AB017563 | Homo sapiens | IGSF4 | 2062 | 99 |
| 208 | X54637 | Homo sapiens | protein tyrosine kinase | 5694 | 98 |
| 209 | AF255910 | Homo sapiens | vascular endothelial junction-associated molecule | 1508 | 98 |
| 210 | AF061324 | Homo sapiens | sulfonylurea receptor 2A | 7545 | 97 |
| 211 | U93568 | Homo sapiens | p40 | 197 | 50 |
| 212 | AF250842 | Drosophila melanogaster | multiple asters | 506 | 32 |
| 213 | X81479 | Homo sapiens | EMR1 | 4469 | 99 |
| 214 | X77748 | Homo sapiens | metabotropic glutamate receptor type 3 (mGIuR3) | 4471 | 99 |
| 215 | M60396 | Homo sapiens | transcobalamin II | 2218 | 99 |
| 216 | W48351 | Homo sapiens | Human breast cancer related protein BCRB2. | 170 | 71 |
| 217 | Y36203 | Homo sapiens | Human secreted protein #75. | 156 | 73 |
| 218 | AF119851 | Homo sapiens | PRO1722 | 144 | 63 |
| 219 | AJ246002 | Mus musculus | spastin protein orthologue | 143 | 100 |
| 220 | D49958 | Homo sapiens | membrane glycoprotein M6 | 616 | 57 |
| 221 | X83573 | Homo sapiens | ARSE | 2114 | 93 |

# TABLE 2

| SEQ ID NO: OF NUCLEOTIDE | ACCESSION NUMBER | SPECIES | DESCRIPTION | SMITH-WATERMAN SCORE | % IDENTITY |
|---|---|---|---|---|---|
| 222 | AF126062 | Homo sapiens | Arf-like 2 binding protein BART1 | 508 | 84 |
| 223 | L22695 | Canine oral papillomavirus | 5' end derived by splicing; putative | 83 | 51 |
| 224 | R95913 | Homo sapiens | Neural thread protein. | 262 | 64 |
| 225 | AP001306 | Arabidopsis thaliana | contains similarity to cell wall-plasma membrane linker protein~gene_id:MKA 23.3 | 79 | 34 |
| 226 | G01984 | Homo sapiens | Human secreted protein, SEQ ID NO: 6065. | 252 | 64 |
| 227 | X04614 | human herpesvirus 1 | IE110 | 83 | 35 |
| 228 | AF151877 | Homo sapiens | CGI-119 protein | 1203 | 94 |
| 229 | AF181467 | Homo sapiens | protein Z-dependent protease inhibitor precursor | 1483 | 88 |
| 230 | Z81326 | Homo sapiens | neuroserpin | 1763 | 99 |
| 231 | AF111173 | Homo sapiens | sodium/hydrogen exchanger isoform 5 | 3512 | 99 |
| 232 | X67055 | Homo sapiens | inter-alpha-trypsin inhibitor heavy chain H3 | 4429 | 98 |
| 233 | AB004064 | Homo sapiens | tomoregulin | 1783 | 98 |
| 234 | AL096772 | Homo sapiens | dJ365O12.1 (KIAA0758 protein) | 5465 | 98 |
| 235 | X83378 | Homo sapiens | putative chloride channel | 1620 | 99 |
| 236 | AF043644 | Homo sapiens | receptor protein tyrosine phosphatase | 5127 | 97 |
| 237 | AF208536 | Homo sapiens | nucleotide binding protein; NBP | 1372 | 100 |
| 238 | AC005625 | Homo sapiens | R27328_1 | 2435 | 93 |
| 239 | X55687 | Lycopersicon esculentum | extensin (class II) | 58 | 50 |
| 240 | M23315 | Sesbania rostrata | nodulin | 61 | 36 |
| 241 | AF102851 | Homo sapiens | dolichyl-P-Glc:Man9GlcNAc2-PP-dolichyl glucosyltransferase | 1881 | 99 |
| 242 | G03793 | Homo sapiens | Human secreted protein, SEQ ID NO: 7874. | 202 | 67 |
| 243 | G03258 | Homo sapiens | Human secreted protein, SEQ ID NO: 7339. | 203 | 69 |
| 244 | AF048774 | Homo sapiens | anti-HER3 scFv | 903 | 81 |

# TABLE 2

| SEQ ID NO: OF NUCLEOTIDE | ACCESSION NUMBER | SPECIES | DESCRIPTION | SMITH-WATERMAN SCORE | % IDENTITY |
|---|---|---|---|---|---|
| 245 | AF102851 | Homo sapiens | dolichyl-P-Glc:Man9GlcNAc2-PP-dolichyl glucosyltransferase | 1867 | 98 |
| 246 | L00352 | Homo sapiens | low density lipoprotein receptor | 3980 | 100 |
| 247 | Y79510 | Homo sapiens | Human carbohydrate-associated protein CRBAP-6. | 1394 | 100 |
| 248 | AF202636 | Homo sapiens | angiopoietin-like protein PP1158 | 2164 | 100 |
| 249 | X66533 | Homo sapiens | guanylate cyclase | 1641 | 97 |
| 250 | M20504 | Homo sapiens | MHC HLA-DR-beta-2 precursor | 750 | 70 |
| 251 | AF157326 | Homo sapiens | TIP120 protein | 4278 | 99 |
| 252 | M25865 | Homo sapiens | von Willebrand factor | 10841 | 95 |
| 253 | AC005625 | Homo sapiens | R27328_1 | 2435 | 93 |
| 254 | A21385 | synthetic construct | heavy chain antibody 3D6 | 1786 | 94 |
| 255 | AF182414 | Homo sapiens | MDS013 | 310 | 48 |
| 256 | Y54041 | Homo sapiens | Protein enc0ded by a gene reduced in metastatic melanoma cells (grmm-1). | 1267 | 84 |
| 257 | AJ011415 | Homo sapiens | plexin-B1/SEP receptor | 1580 | 60 |
| 258 | W55030 | Homo sapiens | G-protein coupled receptor, long form. | 1493 | 100 |
| 259 | AF227747 | Homo sapiens | voltage-dependent calcium channel alpha 1G subunit isoform bc | 6158 | 100 |
| 260 | AF111173 | Homo sapiens | sodium/hydrogen exchanger isoform 5 | 3512 | 99 |
| 261 | G01984 | Homo sapiens | Human secreted protein, SEQ ID NO: 6065. | 175 | 70 |
| 262 | Y00815 | Homo sapiens | put. LAR preprotein (AA -16 to 1881) | 5648 | 100 |
| 263 | Z34979 | Homo sapiens | Human FIZZ3 (inhibitor of neurotrophin action) cDNA. | 582 | 100 |
| 264 | AF119851 | Homo sapiens | PRO1722 | 189 | 73 |
| 265 | AL049798 | Homo sapiens | dJ797M17.1 (Dermatopontin) | 1007 | 99 |
| 266 | AL035684 | Homo sapiens | dJ1114A1.1 (KIAA0611 (putative E1-E2 ATPase) protein) | 1978 | 99 |

# TABLE 2

| SEQ ID NO: OF NUCLEOTIDE | ACCESSION NUMBER | SPECIES | DESCRIPTION | SMITH-WATERMAN SCORE | % IDENTITY |
|---|---|---|---|---|---|
| 267 | U49055 | Rattus norvegicus | rA8 | 4382 | 87 |
| 268 | X15332 | Homo sapiens | alpha-1 (III) collagen | 4170 | 99 |
| 269 | Z98884 | Homo sapiens | dJ467L1.1 (KIAA0833) | 2010 | 100 |
| 270 | AF085244 | Homo sapiens | C2H2 type Kruppel-like zinc finger protein splice variant b | 7331 | 98 |
| 271 | Y00319 | Homo sapiens | Human secreted protein encoded by gene 63. | 214 | 82 |
| 272 | X04434 | Homo sapiens | IGF-I receptor | 5832 | 99 |
| 273 | AC005626 | Homo sapiens | R29124_1 | 1129 | 89 |
| 274 | X52046 | Mus musculus | type III collagen | 819 | 37 |
| 275 | M22207 | Tripneustes gratilla | 217g protein | 168 | 51 |
| 276 | M32317 | Homo sapiens | HLA protein allele B7 | 1536 | 84 |
| 277 | L05485 | Homo sapiens | surfactant protein D | 1693 | 87 |
| 278 | W88504 | Homo sapiens | Human epidermoid carcinoma clone HP10428-encoded membrane protein. | 1187 | 100 |
| 279 | AF078850 | Homo sapiens | steroid dehydrogenase homolog | 794 | 100 |
| 280 | X83378 | Homo sapiens | putative chloride channel | 1620 | 99 |
| 281 | AL035701 | Homo sapiens | dJ8B1.3 (similar to PLASMA-CELL MEMBRANE GLYCOPROTEIN PC-1) | 2412 | 99 |
| 282 | Y87068 | Homo sapiens | Human secreted protein sequence SEQ ID NO:107. | 528 | 100 |
| 283 | L40806 | Neurospora crassa | Restriction enzyme inactivation of met-10 complementation in this region. Sequence similarity to S. cerevisiae chromosome VIII cosmid 9205, accession no. U10556 CDS residues 22627-24126 | 536 | 35 |
| 284 | W88552 | Homo sapiens | Secreted protein encoded by gene 19 clone HSAVU34. | 3078 | 99 |

# TABLE 2

| SEQ ID NO: OF NUCLEOTIDE | ACCESSION NUMBER | SPECIES | DESCRIPTION | SMITH-WATERMAN SCORE | % IDENTITY |
|---|---|---|---|---|---|
| 285 | G03790 | Homo sapiens | Human secreted protein, SEQ ID NO: 7871. | 108 | 50 |
| 286 | X68060 | Homo sapiens | DNA topoisomerase II | 8296 | 99 |
| 287 | G00352 | Homo sapiens | Human secreted protein, SEQ ID NO: 4433. | 114 | 41 |
| 288 | AC004602 | Homo sapiens | F23487_2 | 202 | 49 |
| 289 | AF196329 | Homo sapiens | triggering receptor expressed on monocytes 1 | 1211 | 99 |
| 290 | G03789 | Homo sapiens | Human secreted protein, SEQ ID NO: 7870. | 202 | 62 |
| 291 | G03043 | Homo sapiens | Human secreted protein, SEQ ID NO: 7124. | 93 | 62 |
| 292 | Y12550 | Homo sapiens | Human 5' EST secreted protein SEQ ID NO: 215 from WO 9906553. | 141 | 100 |
| 293 | D43756 | Canis familiaris | fibrinogen A-alpha-chain | 102 | 33 |
| 294 | U38545 | Homo sapiens | phospholipase D1 | 5681 | 99 |
| 295 | W42076 | Homo sapiens | The amino acid sequence of the O276_16 protein. | 236 | 100 |
| 296 | AF090930 | Homo sapiens | PRO0478 | 128 | 60 |
| 297 | Y64747 | Homo sapiens | Human 5' EST related polypeptide SEQ ID NO:908. | 471 | 98 |
| 298 | G01234 | Homo sapiens | Human secreted protein, SEQ ID NO: 5315. | 280 | 71 |
| 299 | G02514 | Homo sapiens | Human secreted protein, SEQ ID NO: 6595. | 94 | 76 |
| 300 | G02493 | Homo sapiens | Human secreted protein, SEQ ID NO: 6574. | 112 | 46 |
| 301 | Z38061 | Saccharomyces cerevisiae | mal5, stal, len: 1367, CAI: 0.3, AMYH_YEAST P08640 GLUCOAMYLASE S1 (EC 3.2.1.3) | 340 | 27 |
| 302 | Y59672 | Homo sapiens | Secreted protein 108-006-5-0-E6-FL. | 530 | 78 |
| 303 | Y95018 | Homo sapiens | Human secreted protein vp19_1, SEQ ID NO:76. | 76 | 35 |
| 304 | W34623 | Homo sapiens | Human C3 protein mutant FT-1. | 117 | 46 |

# TABLE 2

| SEQ ID NO: OF NUCLEOTIDE | ACCESSION NUMBER | SPECIES | DESCRIPTION | SMITH-WATERMAN SCORE | % IDENTITY |
|---|---|---|---|---|---|
| 305 | Y87292 | Homo sapiens | Human signal peptide containing protein HSPP-69 SEQ ID NO:69. | 81 | 50 |
| 306 | AF210651 | Homo sapiens | NAG18 | 135 | 60 |
| 307 | Y14482 | Homo sapiens | Fragment of human secreted protein encoded by gene 17. | 212 | 58 |
| 308 | Y76325 | Homo sapiens | Fragment of human secreted protein encoded by gene 35. | 343 | 93 |
| 309 | Y36156 | Homo sapiens | Human secreted protein #28. | 203 | 75 |
| 310 | AF090931 | Homo sapiens | PRO0483 | 76 | 50 |
| 311 | AC004943 | Homo sapiens | alpha-fetoprotein enhancer-binding protein; 99% identical to A41948 (PID:g283975) | 351 | 85 |
| 312 | G02558 | Homo sapiens | Human secreted protein, SEQ ID NO: 6639. | 144 | 52 |
| 313 | AK000128 | Homo sapiens | unnamed protein product | 1338 | 100 |
| 314 | G03786 | Homo sapiens | Human secreted protein, SEQ ID NO: 7867. | 164 | 83 |
| 315 | AF090942 | Homo sapiens | PRO0657 | 253 | 68 |
| 316 | AF116712 | Homo sapiens | PRO2738 | 181 | 52 |
| 317 | AF043726 | Mus musculus | PHD-finger protein | 1605 | 64 |
| 318 | Y99368 | Homo sapiens | Human PRO1326 (UNQ686) amino acid sequence SEQ ID NO:100. | 145 | 51 |
| 319 | AF065314 | Homo sapiens | cone photoreceptor cGMP-gated channel alpha subunit | 292 | 98 |
| 320 | AF003389 | Caenorhabditis elegans | contains similarity to N-chimaerins | 162 | 28 |
| 321 | Y66755 | Homo sapiens | Membrane-bound protein PRO1185. | 993 | 100 |
| 322 | AF109906 | Mus musculus | RD | 118 | 69 |
| 323 | AF199323 | Rattus norvegicus | RIM2-2A | 364 | 85 |
| 324 | G02538 | Homo sapiens | Human secreted protein, SEQ ID NO: 6619. | 104 | 65 |
| 325 | G02872 | Homo sapiens | Human secreted protein, SEQ ID NO: 6953. | 138 | 65 |
| 326 | Y41266 | Homo sapiens | Human T139 protein. | 591 | 100 |
| 327 | G02920 | Homo sapiens | Human secreted protein, SEQ ID NO: | 103 | 67 |

TABLE 2

| SEQ ID NO: OF NUCLEOTIDE | ACCESSION NUMBER | SPECIES | DESCRIPTION | SMITH-WATERMAN SCORE | % IDENTITY |
|---|---|---|---|---|---|
| | | | 7001. | | |
| 328 | G00636 | Homo sapiens | Human secreted protein, SEQ ID NO: 4717. | 80 | 36 |
| 329 | U37769 | Oryctolagus cuniculus | protein phosphatase 2A0 B' regulatory subunit alpha isoform | 556 | 88 |
| 330 | AE001424 | Plasmodium falciparum | RESA-H3 antigen | 208 | 21 |
| 331 | AF090930 | Homo sapiens | PRO0478 | 156 | 82 |
| 332 | AF161356 | Homo sapiens | HSPC093 | 169 | 64 |
| 333 | G04055 | Homo sapiens | Human secreted protein, SEQ ID NO: 8136. | 425 | 100 |
| 334 | D79985 | Homo sapiens | putative hydrophobic domain in the central region. | 371 | 86 |
| 335 | Y41401 | Homo sapiens | Human secreted protein encoded by gene 94 clone HLYCH68. | 392 | 100 |
| 336 | W18651 | Homo sapiens | Human apolipoprotein E gene +1 frameshift mutant product. | 478 | 88 |
| 337 | Y20921 | Homo sapiens | Human presenilin II wild type protein fragment 5. | 2126 | 96 |
| 338 | AF010144 | Homo sapiens | neuronal thread protein AD7c-NTP | 233 | 75 |
| 339 | D28500 | Homo sapiens | mitochondrial isoleucine tRNA synthetase | 175 | 89 |
| 340 | Y13357 | Homo sapiens | Amino acid sequence of protein PRO227. | 148 | 50 |
| 341 | AL096677 | Homo sapiens | dJ322G13.2 (similar to cystatin) | 94 | 50 |
| 342 | Y10843 | Homo sapiens | Amino acid sequence of a human secreted protein. | 186 | 86 |
| 343 | X54134 | Homo sapiens | protein-tyrosine phosphatase | 3705 | 100 |
| 344 | Z33908 | Mus musculus | inositol 1,4,5-trisphosphate receptor | 315 | 84 |
| 345 | G00241 | Homo sapiens | Human secreted protein, SEQ ID NO: 4322. | 130 | 46 |
| 346 | AF071172 | Homo sapiens | HERC2 | 23705 | 99 |
| 347 | AB015346 | Homo sapiens | Eps15R | 209 | 95 |
| 348 | Y48596 | Homo sapiens | Human breast | 108 | 34 |

# TABLE 2

| SEQ ID NO: OF NUCLEOTIDE | ACCESSION NUMBER | SPECIES | DESCRIPTION | SMITH-WATERMAN SCORE | % IDENTITY |
|---|---|---|---|---|---|
| | | | tumour-associated protein 57. | | |
| 349 | G03058 | Homo sapiens | Human secreted protein, SEQ ID NO: 7139. | 85 | 66 |
| 350 | Y73443 | Homo sapiens | Human secreted protein clone yb187_1 protein sequence SEQ ID NO:108. | 90 | 36 |
| 351 | G03793 | Homo sapiens | Human secreted protein, SEQ ID NO: 7874. | 126 | 66 |
| 352 | G03789 | Homo sapiens | Human secreted protein, SEQ ID NO: 7870. | 324 | 73 |
| 353 | Y64747 | Homo sapiens | Human 5' EST related polypeptide SEQ ID NO:908. | 527 | 98 |
| 354 | AF255342 | Homo sapiens | putative pheromone receptor V1RL1 long form | 147 | 59 |
| 355 | W48351 | Homo sapiens | Human breast cancer related protein BCRB2. | 85 | 61 |
| 356 | G03060 | Homo sapiens | Human secreted protein, SEQ ID NO: 7141. | 191 | 72 |
| 357 | AF124729 | Mus musculus | acinusS' | 124 | 31 |
| 358 | U37352 | Homo sapiens | protein phosphatase 2A B'alpha1 regulatory subunit | 1016 | 95 |
| 359 | AF280605 | Triticum aestivum | omega gliadin storage protein | 125 | 35 |
| 360 | G03789 | Homo sapiens | Human secreted protein, SEQ ID NO: 7870. | 150 | 81 |
| 361 | AL035398 | Homo sapiens | dJ796I17.2 (CGI-51) | 226 | 64 |
| 362 | AK000307 | Homo sapiens | unnamed protein product | 882 | 97 |
| 363 | Y41401 | Homo sapiens | Human secreted protein encoded by gene 94 clone HLYCH68. | 392 | 100 |
| 364 | AF288480 | Homo sapiens | tubby super-family protein | 238 | 87 |
| 365 | AL023706 | Schizosacchar omyces pombe | possible pre-mRNA processing by similarity to yeast prp39 | 383 | 34 |
| 366 | W48351 | Homo sapiens | Human breast cancer related protein BCRB2. | 85 | 61 |

# TABLE 2

| SEQ ID NO: OF NUCLEOTIDE | ACCESSION NUMBER | SPECIES | DESCRIPTION | SMITH-WATERMAN SCORE | % IDENTITY |
|---|---|---|---|---|---|
| 367 | S68978 | Oryctolagus cuniculus | interleukin-1 receptor antagonist intracellular form | 53 | 58 |
| 368 | AF047602 | Equus zebra hartmannae | luteinizing hormone/chorionic gonadotrophin beta-subunit | 68 | 37 |
| 369 | AF119851 | Homo sapiens | PRO1722 | 180 | 75 |
| 370 | U15195 | Homo sapiens | alpha-1 type II collagen | 59 | 43 |
| 371 | U02082 | Homo sapiens | guanine nucleotide regulatory protein | 2648 | 100 |
| 372 | AF096895 | Homo sapiens | chemokine-like factor 1 | 508 | 100 |
| 373 | G03786 | Homo sapiens | Human secreted protein, SEQ ID NO: 7867. | 315 | 65 |
| 374 | AF010144 | Homo sapiens | neuronal thread protein AD7c-NTP | 240 | 67 |
| 375 | U22376 | Homo sapiens | alternatively spliced product using exon 13A | 191 | 80 |
| 376 | U08310 | Saimiri sciureus | prion protein | 245 | 66 |
| 377 | A76867 | unidentified | Chimere G.CSF-Gly4-SAH en aval region prepro de SAH | 550 | 99 |
| 378 | G00442 | Homo sapiens | Human secreted protein, SEQ ID NO: 4523. | 94 | 53 |
| 379 | AF010144 | Homo sapiens | neuronal thread protein AD7c-NTP | 355 | 53 |
| 380 | AB023634 | Rattus norvegicus | Ca/calmodulin-dependent protein kinase phosphatase | 161 | 91 |
| 381 | Y99437 | Homo sapiens | Human PRO1508 (UNQ761) amino acid sequence SEQ ID NO:336. | 805 | 100 |
| 382 | W48351 | Homo sapiens | Human breast cancer related protein BCRB2. | 139 | 61 |
| 383 | M58511 | Homo sapiens | iron-responsive element-binding protein/iron regulatory protein 2 | 286 | 100 |
| 384 | Y02671 | Homo sapiens | Human secreted protein encoded by gene 22 clone HMSJW18. | 99 | 71 |
| 385 | AJ012166 | Canis familiaris | brain-specific synapse associated | 86 | 38 |

# TABLE 2

| SEQ ID NO: OF NUCLEOTIDE | ACCESSION NUMBER | SPECIES | DESCRIPTION | SMITH-WATERMAN SCORE | % IDENTITY |
|---|---|---|---|---|---|
| | | | protein, Bassoon | | |
| 386 | L07809 | Homo sapiens | dynamin | 98 | 31 |
| 387 | M15530 | Homo sapiens | B-cell growth factor | 158 | 69 |
| 388 | AF090172 | Mycoplasma pneumoniae | revertant adhesin-related protein P30 | 109 | 31 |
| 389 | AJ278964 | Homo sapiens | cytosolic beta-glucosidase | 165 | 52 |
| 390 | AF190642 | Homo sapiens | phosphoinositide-specific phospholipase C PLC-epsilon | 1095 | 98 |
| 391 | X13238 | Homo sapiens | cytochrome c oxidase subunit VIc preprotein | 379 | 100 |
| 392 | AF225417 | Homo sapiens | 88.8 kDa protein | 1634 | 98 |
| 393 | Y02693 | Homo sapiens | Human secreted protein encoded by gene 44 clone HTDAD22. | 278 | 75 |
| 394 | AF151037 | Homo sapiens | HSPC203 | 554 | 100 |
| 395 | AJ276396 | Homo sapiens | matrix extracellular phosphoglycoprotein | 465 | 100 |
| 396 | X51405 | Homo sapiens | pre-pro polypeptide (AA -25 to 451) | 2536 | 100 |
| 397 | W78128 | Homo sapiens | Human secreted protein encoded by gene 3 clone HOSBI96. | 564 | 71 |
| 398 | Y87346 | Homo sapiens | Human signal peptide containing protein HSPP-123 SEQ ID NO:123. | 290 | 90 |
| 399 | G03564 | Homo sapiens | Human secreted protein, SEQ ID NO: 7645. | 72 | 52 |
| 400 | U89436 | Homo sapiens | tyrosyl-tRNA synthetase | 2719 | 100 |
| 401 | W80993 | Homo sapiens | Human RIP-interacting factor RIF. | 1724 | 100 |
| 402 | Y27907 | Homo sapiens | Human secreted protein encoded by gene No. 119. | 95 | 59 |
| 403 | AB033102 | Homo sapiens | KIAA1276 protein | 921 | 100 |
| 404 | G03797 | Homo sapiens | Human secreted protein, SEQ ID NO: 7878. | 192 | 55 |
| 405 | AF096895 | Homo sapiens | chemokine-like factor 1 | 508 | 100 |
| 406 | Y29861 | Homo sapiens | Human secreted protein clone | 791 | 98 |

# TABLE 2

| SEQ ID NO: OF NUCLEOTIDE | ACCESSION NUMBER | SPECIES | DESCRIPTION | SMITH-WATERMAN SCORE | % IDENTITY |
|---|---|---|---|---|---|
| | | | cb98_4. | | |
| 407 | Y00293 | Homo sapiens | Human secreted protein encoded by gene 36. | 237 | 97 |
| 408 | W40215 | Homo sapiens | Human macrophage antigen. | 1358 | 99 |
| 409 | L36056 | Homo sapiens | 4E-binding protein 2 | 639 | 100 |
| 410 | AJ130710 | Homo sapiens | QA79 membrane protein, allelic variant airm-1b | 2473 | 100 |
| 411 | AF116661 | Homo sapiens | PRO1438 | 146 | 57 |
| 412 | W88761 | Homo sapiens | Polypeptide fragment encoded by gene 19. | 150 | 58 |
| 413 | AK024434 | Homo sapiens | FLJ00024 protein | 574 | 97 |
| 414 | Y10376 | Homo sapiens | SIRP-beta1 | 2069 | 99 |
| 415 | Y07930 | Homo sapiens | Human secreted protein fragment encoded from gene 79. | 351 | 98 |
| 416 | R99390 | Homo sapiens | Human 030 gene (fohy030) product. | 804 | 71 |
| 417 | AB018253 | Rattus norvegicus | voltage-gated ca channel | 2419 | 88 |
| 418 | AC006017 | Homo sapiens | similar to ALR; similar to AAC51735 (PID:g2358287) | 2150 | 97 |
| 419 | X72925 | Homo sapiens | Dsc1b precursor | 4390 | 99 |
| 420 | AF205940 | Homo sapiens | endomucin | 1289 | 100 |
| 421 | Y27868 | Homo sapiens | Human secreted protein encoded by gene No. 107. | 134 | 54 |
| 422 | W74722 | Homo sapiens | Human secreted protein er80_1. | 2422 | 100 |
| 423 | AF080470 | Homo sapiens | pallid | 872 | 100 |
| 424 | G04072 | Homo sapiens | Human secreted protein, SEQ ID NO: 8153. | 201 | 63 |
| 425 | W90961 | Homo sapiens | Human CSGP-1 protein. | 869 | 86 |
| 426 | M13180 | Human herpesvirus 4 | nuclear antigen (EBNA 1) | 59 | 45 |
| 427 | G00365 | Homo sapiens | Human secreted protein, SEQ ID NO: 4446. | 99 | 75 |
| 428 | AF155819 | Mus musculus | doublecortin-like kinase | 3448 | 96 |
| 429 | Y04315 | Homo sapiens | Human secreted protein encoded by gene 23. | 385 | 100 |
| 430 | AB026891 | Homo sapiens | cystine/glutamate transporter | 2552 | 100 |

# TABLE 2

| SEQ ID NO: OF NUCLEOTIDE | ACCESSION NUMBER | SPECIES | DESCRIPTION | SMITH-WATERMAN SCORE | % IDENTITY |
|---|---|---|---|---|---|
| 431 | Y15286 | Homo sapiens | vacuolar proton-ATPase subunit M9.2 | 459 | 100 |
| 432 | X81053 | Homo sapiens | type IV collagen alpha 4 chain | 9706 | 99 |
| 433 | U41829 | Macaca mulatta | MHC class I antigen Mamu B*07 | 365 | 76 |
| 434 | G03371 | Homo sapiens | Human secreted protein, SEQ ID NO: 7452. | 100 | 41 |
| 435 | AF233238 | Gallus gallus | BMP signal transducer Smad1 | 170 | 74 |
| 436 | X52425 | Homo sapiens | interleukin 4 receptor | 4492 | 99 |
| 437 | Y06115 | Homo sapiens | Human organic cation transporter OCT-3. | 2593 | 96 |
| 438 | G02872 | Homo sapiens | Human secreted protein, SEQ ID NO: 6953. | 130 | 54 |
| 439 | L08239 | Homo sapiens | located at OATL1 | 1304 | 95 |
| 440 | X17115 | Homo sapiens | precursor (AA -15 to 612) | 2613 | 86 |
| 441 | Y06816 | Homo sapiens | Human Notch2 (humN2) protein sequence. | 1471 | 98 |
| 442 | AB019440 | Homo sapiens | immunogloblin heavy chain variable region | 545 | 88 |
| 443 | Y87350 | Homo sapiens | Human signal peptide containing protein HSPP-127 SEQ ID NO:127. | 1061 | 100 |
| 444 | AJ271736 | Homo sapiens | synaptobrevin-like 1 protein | 1128 | 100 |
| 445 | Y11534 | Homo sapiens | PEG1/MEST | 1787 | 100 |
| 446 | W85719 | Homo sapiens | Novel protein (Clone AJ143_1). | 271 | 100 |
| 447 | Y07900 | Homo sapiens | Human secreted protein fragment encoded from gene 49. | 87 | 94 |
| 448 | X14329 | Homo sapiens | carboxypeptidase N precursor (AA -20 to 438) | 2463 | 99 |
| 449 | M36803 | Homo sapiens | hemopexin | 2603 | 100 |
| 450 | AF116238 | Homo sapiens | pseudouridine synthase 1 | 1927 | 99 |
| 451 | AB031051 | Homo sapiens | organic anion transporter OATP-E | 444 | 42 |
| 452 | X16841 | Homo sapiens | precursor protein (-19 to 742) | 3958 | 100 |
| 453 | AK022830 | Homo sapiens | unnamed protein product | 373 | 100 |

# TABLE 2

| SEQ ID NO: OF NUCLEOTIDE | ACCESSION NUMBER | SPECIES | DESCRIPTION | SMITH-WATERMAN SCORE | % IDENTITY |
|---|---|---|---|---|---|
| 454 | Y94890 | Homo sapiens | Human protein clone HP02798. | 637 | 90 |
| 455 | AL356014 | Arabidopsis thaliana | putative protein | 210 | 38 |
| 456 | X60221 | Homo sapiens | H+-ATP synthase subunit b | 1297 | 99 |
| 457 | G02532 | Homo sapiens | Human secreted protein, SEQ ID NO: 6613. | 168 | 69 |
| 458 | AJ245375 | Homo sapiens | PP35 act | 1895 | 99 |
| 459 | G00397 | Homo sapiens | Human secreted protein, SEQ ID NO: 4478. | 57 | 52 |
| 460 | AE003708 | Drosophila melanogaster | CG6194 gene product | 234 | 65 |
| 461 | W48352 | Homo sapiens | Human breast cancer related protein BCFLT1. | 80 | 60 |
| 462 | U53420 | Rattus norvegicus | sodium-calcium exchanger form 3 | 397 | 76 |
| 463 | Y13402 | Homo sapiens | Amino acid sequence of protein PRO310. | 1075 | 63 |
| 464 | Y27607 | Homo sapiens | Human secreted protein encoded by gene No. 41. | 610 | 100 |
| 465 | L08666 | Homo sapiens | porin | 122 | 51 |
| 466 | Y87084 | Homo sapiens | Human secreted protein sequence SEQ ID NO:123. | 232 | 78 |
| 467 | X16841 | Homo sapiens | precursor protein (-19 to 742) | 3958 | 100 |
| 468 | Y48507 | Homo sapiens | Human breast tumour-associated protein 52. | 295 | 91 |
| 469 | X07973 | Ovis aries | MT-Ib protein | 84 | 45 |
| 470 | W48927 | Homo sapiens | Schwannomin-binding protein C-terminal fragment. | 78 | 60 |
| 471 | AJ224171 | Homo sapiens | lipophilin A | 454 | 100 |
| 472 | G01984 | Homo sapiens | Human secreted protein, SEQ ID NO: 6065. | 211 | 64 |
| 473 | G03793 | Homo sapiens | Human secreted protein, SEQ ID NO: 7874. | 200 | 74 |
| 474 | Y17829 | Homo sapiens | Human PRO354 protein sequence. | 1006 | 100 |
| 475 | Y66706 | Homo sapiens | Membrane-bound protein PRO1129. | 2153 | 99 |
| 476 | G03800 | Homo sapiens | Human secreted protein, SEQ ID NO: 7881. | 99 | 78 |
| 477 | AF216389 | Homo sapiens | semaphorin Rs | 296 | 85 |

# TABLE 2

| SEQ ID NO: OF NUCLEOTIDE | ACCESSION NUMBER | SPECIES | DESCRIPTION | SMITH-WATERMAN SCORE | % IDENTITY |
|---|---|---|---|---|---|
| 478 | X93036 | Homo sapiens | MAT8 protein | 469 | 100 |
| 479 | X53795 | Homo sapiens | inducible membrane protein | 1412 | 100 |
| 480 | AF056195 | Homo sapiens | neuroblastoma-amplified protein | 4504 | 98 |
| 481 | AF116715 | Homo sapiens | PRO2829 | 96 | 46 |
| 482 | Z24680 | Homo sapiens | garp | 167 | 43 |
| 483 | Y76198 | Homo sapiens | Human secreted protein encoded by gene 75. | 82 | 80 |
| 484 | AF010144 | Homo sapiens | neuronal thread protein AD7c-NTP | 324 | 59 |
| 485 | Y91592 | Homo sapiens | Human secreted protein sequence encoded by gene 6 SEQ ID NO:265. | 738 | 100 |
| 486 | Y94890 | Homo sapiens | Human protein clone HP02798. | 605 | 81 |
| 487 | U89436 | Homo sapiens | tyrosyl-tRNA synthetase | 2719 | 100 |
| 488 | W88579 | Homo sapiens | Secreted protein encoded by gene 46 clone HCFMV39. | 479 | 95 |
| 489 | G02360 | Homo sapiens | Human secreted protein, SEQ ID NO: 6441. | 102 | 70 |
| 490 | U70976 | Homo sapiens | arrestin | 1071 | 61 |
| 491 | U80746 | Homo sapiens | CAGH4 | 277 | 81 |
| 492 | U26361 | Helicobacter pylori | Hpn | 80 | 83 |
| 493 | Y19730 | Homo sapiens | SEQ ID NO 448 from WO9922243. | 135 | 53 |
| 494 | Y27868 | Homo sapiens | Human secreted protein encoded by gene No. 107. | 185 | 50 |
| 495 | AF090901 | Homo sapiens | PRO0195 | 90 | 46 |
| 496 | AF061529 | Mus musculus | rjs | 270 | 76 |
| 497 | L34049 | Rattus norvegicus | megalin | 322 | 41 |
| 498 | J04204 | Bos taurus | 32 kd accessory protein | 1743 | 100 |
| 499 | Y71118 | Homo sapiens | Human Hydrolase protein-16 (HYDRL-16). | 2205 | 97 |
| 500 | X13916 | Homo sapiens | LDL-receptor related precursor (AA -19 to 4525) | 715 | 92 |
| 501 | Y00877 | Homo sapiens | Human LAPH-2 protein sequence. | 138 | 40 |
| 502 | Y99368 | Homo sapiens | Human PRO1326 (UNQ686) amino acid sequence SEQ ID NO:100. | 156 | 48 |

# TABLE 2

| SEQ ID NO: OF NUCLEOTIDE | ACCESSION NUMBER | SPECIES | DESCRIPTION | SMITH-WATERMAN SCORE | % IDENTITY |
|---|---|---|---|---|---|
| 503 | Y48308 | Homo sapiens | Human prostate cancer-associated protein 5. | 901 | 100 |
| 504 | U67060 | Cricetulus griseus | SREBP cleavage activating protein | 6196 | 92 |
| 505 | W75857 | Homo sapiens | Human secretory protein of clone CO1020-1. | 1761 | 99 |
| 506 | X55764 | Homo sapiens | 11beta-hydrolase precursor | 2604 | 99 |
| 507 | Y41685 | Homo sapiens | Human PRO213 protein sequence. | 1344 | 94 |
| 508 | X95240 | Homo sapiens | cysteine-rich secretory protein-3 | 1368 | 100 |
| 509 | AF065482 | Homo sapiens | sorting nexin 2 | 517 | 77 |
| 510 | AF135025 | Homo sapiens | kallikrein-like protein 5-related protein 1 | 1301 | 100 |
| 511 | AF220492 | Homo sapiens | krueppel-like zinc finger protein HZF2 | 4100 | 99 |
| 512 | X58397 | Homo sapiens | variable region V251 from V(H)5 gene | 670 | 100 |
| 513 | W95348 | Homo sapiens | Human foetal kidney secreted protein em397_2. | 406 | 90 |
| 514 | AJ000479 | Homo sapiens | putative G-Protein coupled receptor, EDG6 | 1966 | 100 |
| 515 | L05514 | Homo sapiens | histatin 3 | 280 | 100 |
| 516 | X95240 | Homo sapiens | cysteine-rich secretory protein-3 | 1368 | 100 |
| 517 | D00654 | Homo sapiens | enteric smooth muscle gamma-actin | 1972 | 100 |
| 518 | AJ005453 | Mytilus edulis | metallothionein 10 II | 94 | 35 |
| 519 | W37864 | Homo sapiens | Human protein comprising secretory signal amino acid sequence 1. | 362 | 98 |
| 520 | X76091 | Homo sapiens | DNA binding protein RFX2 | 3743 | 99 |
| 521 | G03800 | Homo sapiens | Human secreted protein, SEQ ID NO: 7881. | 113 | 39 |
| 522 | AJ289243 | Mus musculus | calpain 12 | 147 | 53 |
| 523 | D30037 | Homo sapiens | phosphatidylinositol transfer protein | 1464 | 100 |
| 524 | AJ012370 | Homo sapiens | NAALADase II protein | 3872 | 99 |
| 525 | G03909 | Homo sapiens | Human secreted protein, SEQ ID NO: | 80 | 41 |

# TABLE 2

| SEQ ID NO: OF NUCLEOTIDE | ACCESSION NUMBER | SPECIES | DESCRIPTION | SMITH-WATERMAN SCORE | % IDENTITY |
|---|---|---|---|---|---|
| | | | 7990. | | |
| 526 | U67060 | Cricetulus griseus | SREBP cleavage activating protein | 6196 | 92 |
| 527 | W48351 | Homo sapiens | Human breast cancer related protein BCRB2. | 85 | 61 |
| 528 | AF093408 | Homo sapiens | protein kinase A binding protein AKAP110 | 461 | 78 |
| 529 | Y92182 | Homo sapiens | Human partial TANGO 195 from clone T195Athpb93f1. | 1682 | 100 |
| 530 | M28200 | Homo sapiens | MHC class II lymphocyte antigen beta chain | 432 | 72 |
| 531 | X58397 | Homo sapiens | variable region V251 from V(H)5 gene | 491 | 74 |
| 532 | D88577 | Mus musculus | Kupffer cell receptor | 904 | 46 |
| 533 | M84379 | Homo sapiens | lymphocyte antigen | 1922 | 97 |
| 534 | AF279265 | Homo sapiens | putative anion transporter 1 | 212 | 91 |
| 535 | AF132035 | Homo sapiens | core 2 beta-1,6-N-acetylglucosaminyltransferase 3 | 852 | 92 |
| 536 | G02958 | Homo sapiens | Human secreted protein, SEQ ID NO: 7039. | 512 | 98 |
| 537 | Y07938 | Homo sapiens | Human secreted protein fragment encoded from gene 87. | 302 | 100 |
| 538 | Y36203 | Homo sapiens | Human secreted protein #75. | 175 | 51 |
| 539 | U16738 | Homo sapiens | CAG-isl 7 | 472 | 75 |
| 540 | AL161531 | Arabidopsis thaliana | putative proline-rich protein | 118 | 57 |
| 541 | K00558 | Homo sapiens | alpha-tubulin | 2393 | 100 |
| 542 | U20286 | Rattus norvegicus | lamina associated polypeptide 1C | 641 | 55 |
| 543 | Y27907 | Homo sapiens | Human secreted protein encoded by gene No. 119. | 128 | 61 |
| 544 | AF109674 | Rattus norvegicus | late gestation lung protein 1 | 954 | 87 |
| 545 | L35278 | Homo sapiens | bone morphogenetic protein | 92 | 40 |
| 546 | G00541 | Homo sapiens | Human secreted protein, SEQ ID NO: 4622. | 94 | 68 |
| 547 | AF190664 | Mus musculus | LMBR2 | 246 | 78 |
| 548 | Y12793 | Homo sapiens | Human 5' EST | 113 | 50 |

# TABLE 2

| SEQ ID NO: OF NUCLEOTIDE | ACCESSION NUMBER | SPECIES | DESCRIPTION | SMITH-WATERMAN SCORE | % IDENTITY |
|---|---|---|---|---|---|
| | | | secreted protein SEQ ID NO:383. | | |
| 549 | AF133816 | Homo sapiens | insulin-like peptide INSL5 | 714 | 100 |
| 550 | X70910 | Homo sapiens | tetranectin | 1069 | 100 |
| 551 | M11902 | Mus musculus | proline-rich salivary protein | 135 | 39 |
| 552 | G03477 | Homo sapiens | Human secreted protein, SEQ ID NO: 7558. | 89 | 58 |
| 553 | U63542 | Homo sapiens | FAP protein | 156 | 77 |
| 554 | Y60497 | Homo sapiens | Human normal bladder tissue EST encoded protein 169. | 89 | 50 |
| 555 | Y87303 | Homo sapiens | Human signal peptide containing protein HSPP-80 SEQ ID NO:80. | 275 | 100 |
| 556 | Y17526 | Homo sapiens | Human secreted protein clone AM349 2 protein. | 1220 | 100 |
| 557 | G04064 | Homo sapiens | Human secreted protein, SEQ ID NO: 8145. | 83 | 35 |
| 558 | U51919 | Rattus norvegicus | preprocortistatin | 84 | 36 |
| 559 | AF090901 | Homo sapiens | PRO0195 | 92 | 66 |
| 560 | J04031 | Homo sapiens | MDMCSF (EC 1.5.1.5; EC 3.5.4.9; EC 6.3.4.3) | 226 | 52 |
| 561 | AL117237 | Homo sapiens | hypothetical protein | 4088 | 94 |
| 562 | Y50931 | Homo sapiens | Human fetal brain cDNA clone vc25_1 derived protein. | 485 | 100 |
| 563 | Y21631 | Homo sapiens | Ligand binding domain of nuclear receptor hTRbeta. | 1738 | 99 |
| 564 | X90857 | Homo sapiens | -14 | 177 | 69 |
| 565 | W35904 | Homo sapiens | Human haematopoietic-specific protein (HSP). | 862 | 87 |
| 566 | W99070 | Homo sapiens | Human PIGR-1. | 244 | 90 |
| 567 | X61653 | Homo sapiens | TCR V-beta 13.5 | 600 | 100 |
| 568 | AF166350 | Homo sapiens | ST7 protein | 4711 | 99 |
| 569 | Y07938 | Homo sapiens | Human secreted protein fragment encoded from gene 87. | 302 | 100 |
| 570 | X85019 | Homo sapiens | UDP-GalNAc:polypeptide | 3069 | 100 |

TABLE 2

| SEQ ID NO: OF NUCLEOTIDE | ACCESSION NUMBER | SPECIES | DESCRIPTION | SMITH-WATERMAN SCORE | % IDENTITY |
|---|---|---|---|---|---|
| | | | N-acetylgalactosaminyl transferase | | |
| 571 | U89942 | Homo sapiens | lysyl oxidase-related protein | 2427 | 89 |
| 572 | X04391 | Homo sapiens | put. precursor polypeptide | 2671 | 99 |
| 573 | W36903 | Homo sapiens | Human epididymis-specific receptor protein. | 5352 | 100 |
| 574 | U22816 | Homo sapiens | LAR-interacting protein 1b | 2042 | 57 |
| 575 | Y58618 | Homo sapiens | Protein regulating gene expression PRGE-11. | 729 | 57 |
| 576 | AJ278348 | Homo sapiens | pregnancy-associated plasma protein-E | 743 | 100 |
| 577 | AK024512 | Homo sapiens | unnamed protein product | 471 | 100 |
| 578 | AL031685 | Homo sapiens | dJ963K23.4 (KIAA0939 (novel Sodium/hydrogen exchanger family member)) | 2010 | 100 |
| 579 | AF183183 | Mus musculus | cochlear otoferlin | 116 | 91 |
| 580 | W74722 | Homo sapiens | Human secreted protein er80_1. | 2422 | 100 |
| 581 | G03356 | Homo sapiens | Human secreted protein, SEQ ID NO: 7437. | 114 | 44 |
| 582 | Y82777 | Homo sapiens | Human chordin related protein (Clone dw665_4). | 610 | 98 |
| 583 | J04988 | Homo sapiens | 90 kD heat shock protein | 3702 | 100 |
| 584 | K02576 | Homo sapiens | salivary proline-rich protein 1 | 97 | 34 |
| 585 | G03786 | Homo sapiens | Human secreted protein, SEQ ID NO: 7867. | 159 | 72 |
| 586 | AK024490 | Homo sapiens | FLJ00092 protein | 204 | 57 |
| 587 | U22231 | Felis catus | ribosomal protein S3a | 327 | 57 |
| 588 | X55681 | Lycopersicon esculentum | extensin (class I) | 96 | 38 |
| 589 | U68137 | Rana ridibunda | prepro-somatostatin 14 | 81 | 33 |
| 590 | Y19655 | Homo sapiens | SEQ ID NO 373 from WO9922243. | 814 | 84 |
| 591 | G03789 | Homo sapiens | Human secreted protein, SEQ ID NO: 7870. | 222 | 56 |

# TABLE 2

| SEQ ID NO: OF NUCLEOTIDE | ACCESSION NUMBER | SPECIES | DESCRIPTION | SMITH-WATERMAN SCORE | % IDENTITY |
|---|---|---|---|---|---|
| 592 | AF067801 | Homo sapiens | HDCGC21P | 116 | 38 |
| 593 | X67339 | Neurospora crassa | ccg-2 | 82 | 37 |
| 594 | G03280 | Homo sapiens | Human secreted protein, SEQ ID NO: 7361. | 169 | 100 |
| 595 | Y02693 | Homo sapiens | Human secreted protein encoded by gene 44 clone HTDAD22. | 130 | 70 |
| 596 | AE003683 | Drosophila melanogaster | CG9492 gene product | 247 | 56 |
| 597 | Z22968 | Homo sapiens | M130 antigen | 6205 | 100 |
| 598 | AK021847 | Homo sapiens | unnamed protein product | 178 | 94 |
| 599 | AP000060 | Aeropyrum pernix | 134aa long hypothetical protein | 80 | 39 |
| 600 | AK001363 | Homo sapiens | unnamed protein product | 558 | 92 |
| 601 | G02872 | Homo sapiens | Human secreted protein, SEQ ID NO: 6953. | 147 | 49 |
| 602 | G02538 | Homo sapiens | Human secreted protein, SEQ ID NO: 6619. | 149 | 65 |
| 603 | X98330 | Homo sapiens | ryanodine receptor 2 | 25918 | 99 |
| 604 | AJ243460 | Leishmania major | proteophosphoglycan | 172 | 35 |
| 605 | Y81807 | Homo sapiens | Human mahogany protein sequence #2. | 2499 | 63 |
| 606 | AF041069 | Equus caballus | fibronectin | 109 | 56 |
| 607 | Y54591 | Homo sapiens | Amino acid sequence of a human transferase designated HUTRAN-1. | 153 | 77 |
| 608 | G03172 | Homo sapiens | Human secreted protein, SEQ ID NO: 7253. | 82 | 66 |
| 609 | Y31730 | Homo sapiens | Human fused protein kinase-deletion mutant fused C-term. | 561 | 99 |
| 610 | Y30163 | Homo sapiens | Human dorsal root receptor 5 hDRR5. | 112 | 49 |
| 611 | G03714 | Homo sapiens | Human secreted protein, SEQ ID NO: 7795. | 171 | 70 |
| 612 | U58514 | Homo sapiens | chitinase precursor | 402 | 75 |

# TABLE 2

| SEQ ID NO: OF NUCLEOTIDE | ACCESSION NUMBER | SPECIES | DESCRIPTION | SMITH-WATERMAN SCORE | % IDENTITY |
|---|---|---|---|---|---|
| 613 | AL122105 | Homo sapiens | hypothetical protein | 399 | 73 |
| 614 | AF059198 | Homo sapiens | protein kinase/endoribonulcease | 5093 | 99 |
| 615 | X17531 | Strongylocentrotus purpuratus | epidermal growth factor | 234 | 54 |
| 616 | AF112982 | Homo sapiens | group IID secretory phospholipase A2 | 852 | 100 |
| 617 | AJ006119 | Homo sapiens | anti-IFN-G scFv | 675 | 97 |
| 618 | W54097 | Homo sapiens | Homo sapiens B223 sequence. | 339 | 98 |
| 619 | AF090930 | Homo sapiens | PRO0478 | 141 | 79 |
| 620 | W61624 | Homo sapiens | Clone HHFEK40 of TM4SF superfamily. | 564 | 98 |
| 621 | AF119851 | Homo sapiens | PRO1722 | 115 | 52 |
| 622 | G03172 | Homo sapiens | Human secreted protein, SEQ ID NO: 7253. | 173 | 48 |
| 623 | Y41379 | Homo sapiens | Human secreted protein encoded by gene 72 clone HE6GA29. | 261 | 100 |
| 624 | U86339 | Drosophila grimshawi | expanded | 142 | 36 |
| 625 | D86853 | Catharanthus roseus | extensin | 142 | 39 |
| 626 | S58722 | Homo sapiens | X-linked retinopathy protein {C-terminal, clone XEH.8c} | 116 | 49 |
| 627 | G02532 | Homo sapiens | Human secreted protein, SEQ ID NO: 6613. | 108 | 50 |
| 628 | G03790 | Homo sapiens | Human secreted protein, SEQ ID NO: 7871. | 129 | 61 |
| 629 | Y27665 | Homo sapiens | Human secreted protein encoded by gene No. 99. | 345 | 100 |
| 630 | G02837 | Homo sapiens | Human secreted protein, SEQ ID NO: 6918. | 78 | 75 |
| 631 | G03789 | Homo sapiens | Human secreted protein, SEQ ID NO: 7870. | 172 | 65 |
| 632 | X14329 | Homo sapiens | carboxypeptidase N precursor (AA -20 to 438) | 2463 | 99 |
| 633 | Y87235 | Homo sapiens | Human signal peptide containing protein HSPP-12 SEQ | 867 | 100 |

# TABLE 2

| SEQ ID NO: OF NUCLEOTIDE | ACCESSION NUMBER | SPECIES | DESCRIPTION | SMITH-WATERMAN SCORE | % IDENTITY |
|---|---|---|---|---|---|
| | | | ID NO:12. | | |
| 634 | W88627 | Homo sapiens | Secreted protein encoded by gene 94 clone HPMBQ32. | 106 | 73 |
| 635 | W74845 | Homo sapiens | Human secreted protein encoded by gene 117 clone HBMUW78. | 395 | 71 |
| 636 | M16941 | Homo sapiens | DR7 beta-chain glycoprotein | 1412 | 100 |
| 637 | W95634 | Homo sapiens | Homo sapiens secreted protein. | 1391 | 100 |
| 638 | Y78801 | Homo sapiens | Hydrophobic domain containing protein clone HP00631 amino acid sequence. | 1277 | 100 |
| 639 | G03789 | Homo sapiens | Human secreted protein, SEQ ID NO: 7870. | 191 | 76 |
| 640 | W64535 | Homo sapiens | Human leukocyte cell clone HP00804 protein. | 2014 | 99 |
| 641 | Y94621 | Homo sapiens | Epidermal growth factor-like variant in skin-2 amino acid sequence. | 529 | 91 |
| 642 | G03646 | Homo sapiens | Human secreted protein, SEQ ID NO: 7727. | 81 | 42 |
| 643 | Y87328 | Homo sapiens | Human signal peptide containing protein HSPP-105 SEQ ID NO:105. | 681 | 100 |
| 644 | Y21386 | Homo sapiens | Human HUPF-I mutant protein fragment 34. | 78 | 31 |
| 645 | G03790 | Homo sapiens | Human secreted protein, SEQ ID NO: 7871. | 140 | 55 |
| 646 | Y35894 | Homo sapiens | Extended human secreted protein sequence, SEQ ID NO. 143. | 349 | 100 |
| 647 | G00517 | Homo sapiens | Human secreted protein, SEQ ID NO: 4598. | 109 | 37 |
| 648 | Y25716 | Homo sapiens | Human secreted protein encoded from gene 6. | 339 | 39 |
| 649 | G01246 | Homo sapiens | Human secreted protein, SEQ ID NO: 5327. | 152 | 80 |
| 650 | R95913 | Homo sapiens | Neural thread | 233 | 50 |

## TABLE 2

| SEQ ID NO: OF NUCLEOTIDE | ACCESSION NUMBER | SPECIES | DESCRIPTION | SMITH-WATERMAN SCORE | % IDENTITY |
|---|---|---|---|---|---|
| | | | protein. | | |
| 651 | Y91469 | Homo sapiens | Human secreted protein sequence encoded by gene 19 SEQ ID NO:142. | 98 | 48 |
| 652 | G03136 | Homo sapiens | Human secreted protein, SEQ ID NO: 7217. | 94 | 43 |
| 653 | U14635 | Caenorhabditis elegans | weak similarity to NADH dehydrogenase | 186 | 30 |
| 654 | Y14482 | Homo sapiens | Fragment of human secreted protein encoded by gene 17. | 163 | 54 |
| 655 | U14635 | Caenorhabditis elegans | weak similarity to NADH dehydrogenase | 186 | 30 |
| 656 | AB024565 | Mus musculus | heparan sulfate 6-sulfotransferase 2 | 1128 | 79 |
| 657 | G03789 | Homo sapiens | Human secreted protein, SEQ ID NO: 7870. | 243 | 70 |
| 658 | Y14471 | Homo sapiens | Fragment of human secreted protein encoded by gene 4. | 95 | 65 |
| 659 | AF135381 | Homo sapiens | chemokine-like factor 3 | 89 | 59 |
| 660 | U40407 | synthetic construct | T cell receptor alpha chain | 586 | 100 |
| 661 | AF039712 | Caenorhabditis elegans | contains similarity to CDP-alcohol phosphotransferases | 289 | 43 |
| 662 | G03790 | Homo sapiens | Human secreted protein, SEQ ID NO: 7871. | 113 | 55 |
| 663 | AF084467 | Homo sapiens | heparanase | 170 | 32 |
| 664 | AF279890 | Homo sapiens | 2P domain potassium channel TREK2 | 1189 | 94 |
| 665 | W63693 | Homo sapiens | Human secreted protein 13. | 243 | 84 |
| 666 | AE003908 | Xylella fastidiosa | hypothetical protein | 120 | 28 |
| 667 | B08948 | Homo sapiens | Human secreted protein sequence encoded by gene 21 SEQ ID NO:105. | 985 | 89 |
| 668 | AF023158 | Homo sapiens | tyrosine phosphatase | 346 | 64 |
| 669 | AF169257 | Homo sapiens | sodium/calcium exchanger NCKX3 | 189 | 57 |
| 670 | AF132969 | Homo sapiens | CGI-35 protein | 364 | 69 |
| 671 | AF269286 | Homo sapiens | HC6 | 112 | 50 |
| 672 | X98494 | Homo sapiens | M phase phosphoprotein 10 | 529 | 68 |
| 673 | G03787 | Homo sapiens | Human secreted | 83 | 44 |

# TABLE 2

| SEQ ID NO: OF NUCLEOTIDE | ACCESSION NUMBER | SPECIES | DESCRIPTION | SMITH-WATERMAN SCORE | % IDENTITY |
|---|---|---|---|---|---|
| | | | protein, SEQ ID NO: 7868. | | |
| 674 | AF119855 | Homo sapiens | PRO1847 | 123 | 46 |
| 675 | AJ242540 | Volvox carteri f. nagariensis | hydroxyproline-rich glycoprotein DZ-HRGP | 242 | 42 |
| 676 | Y91666 | Homo sapiens | Human secreted protein sequence encoded by gene 72 SEQ ID NO:339. | 529 | 96 |
| 677 | Y57936 | Homo sapiens | Human transmembrane protein HTMPN-60. | 669 | 100 |
| 678 | G03789 | Homo sapiens | Human secreted protein, SEQ ID NO: 7870. | 156 | 72 |
| 679 | W18878 | Homo sapiens | Human protein kinase C inhibitor, IPKC-1. | 98 | 68 |
| 680 | Z12168 | Canis familiaris | stimulatory GTP binding protein | 980 | 88 |
| 681 | G00517 | Homo sapiens | Human secreted protein, SEQ ID NO: 4598. | 160 | 48 |
| 682 | W19932 | Homo sapiens | Alzheimer's disease protein encoded by DNA from plasmid pGCS55. | 362 | 100 |
| 683 | Y30709 | Homo sapiens | Amino acid sequence of a human secreted protein. | 99 | 56 |
| 684 | AF269286 | Homo sapiens | HC6 | 137 | 72 |
| 685 | M14362 | Homo sapiens | T-cell surface antigen CD2 precursor | 275 | 64 |
| 686 | G02493 | Homo sapiens | Human secreted protein, SEQ ID NO: 6574. | 173 | 61 |
| 687 | AF248635 | Mus musculus | lymphocyte antigen 108 isoform 1 | 303 | 50 |
| 688 | D86983 | Homo sapiens | similar to D.melanogaster peroxidasin(U11052) | 288 | 55 |
| 689 | Y59711 | Homo sapiens | Secreted protein 58-20-4-G7-FL1. | 895 | 91 |
| 690 | W48848 | Homo sapiens | Human receptor tyrosine kinase LMR3_h N-terminal polypeptide. | 1056 | 89 |
| 691 | W22652 | Homo sapiens | 64-863 antibody HSV863 light chain variable region. | 459 | 77 |
| 692 | AF098066 | Homo sapiens | squamous cell carcinoma antigen | 1001 | 98 |

# TABLE 2

| SEQ ID NO: OF NUCLEOTIDE | ACCESSION NUMBER | SPECIES | DESCRIPTION | SMITH-WATERMAN SCORE | % IDENTITY |
|---|---|---|---|---|---|
| | | | recognized by T cell | | |
| 693 | D83039 | Homo sapiens | eti-1 | 426 | 98 |
| 694 | Y79511 | Homo sapiens | Human carbohydrate-associated protein CRBAP-7. | 1245 | 99 |
| 695 | U12623 | Rattus norvegicus | cyclic nucleotide gated cation channel | 857 | 83 |
| 696 | AF229067 | Homo sapiens | PADI-H protein | 174 | 61 |
| 697 | G03789 | Homo sapiens | Human secreted protein, SEQ ID NO: 7870. | 196 | 75 |
| 698 | U10921 | Macaca mulatta | T-cell receptor alpha chain | 578 | 82 |
| 699 | U31913 | Homo sapiens | HBV-X associated protein | 167 | 100 |
| 700 | X99043 | Mus musculus | brain-derived immunoglobulin superfamily molecule | 348 | 82 |
| 701 | X59770 | Homo sapiens | type II interleukin-1 receptor | 2130 | 100 |
| 702 | AC018758 | Homo sapiens | GPI-anchored metastasis-associated protein homolog | 207 | 31 |
| 703 | Y28816 | Homo sapiens | pm4_13 secreted protein. | 280 | 100 |
| 704 | Y52386 | Homo sapiens | Human transmembrane protein HP02000. | 1077 | 100 |
| 705 | U12392 | Haematobia irritans | putative ATPase | 481 | 55 |
| 706 | U11265 | Homo sapiens | HLA-B35 | 351 | 92 |
| 707 | X64594 | Homo sapiens | 50 kDa erythrocyte plasma membrane glycoprotein | 301 | 88 |
| 708 | AB046048 | Macaca fascicularis | unnamed portein product | 260 | 67 |
| 709 | G03807 | Homo sapiens | Human secreted protein, SEQ ID NO: 7888. | 119 | 60 |
| 710 | G03315 | Homo sapiens | Human secreted protein, SEQ ID NO: 7396. | 314 | 100 |
| 711 | Y50945 | Homo sapiens | Human adult thymus cDNA clone vh1_1 derived protein #1. | 742 | 100 |
| 712 | G00564 | Homo sapiens | Human secreted protein, SEQ ID NO: 4645. | 271 | 98 |
| 713 | G00125 | Homo sapiens | Human secreted | 373 | 80 |

# TABLE 2

| SEQ ID NO: OF NUCLEOTIDE | ACCESSION NUMBER | SPECIES | DESCRIPTION | SMITH-WATERMAN SCORE | % IDENTITY |
|---|---|---|---|---|---|
| | | | protein, SEQ ID NO: 4206. | | |
| 714 | Y13352 | Homo sapiens | Amino acid sequence of protein PRO228. | 872 | 98 |
| 715 | G02753 | Homo sapiens | Human secreted protein, SEQ ID NO: 6834. | 222 | 68 |
| 716 | Y19588 | Homo sapiens | Amino acid sequence of a human secreted protein. | 329 | 100 |
| 717 | AB030235 | Canis familiaris | D4 dopamine receptor | 79 | 35 |
| 718 | W74577 | Homo sapiens | Human membrane protein BA2303. | 748 | 100 |
| 719 | Y02693 | Homo sapiens | Human secreted protein encoded by gene 44 clone HTDAD22. | 235 | 61 |
| 720 | X97868 | Homo sapiens | arylsulphatase | 167 | 84 |
| 721 | Y13215 | Homo sapiens | Human secreted protein encoded by 5' EST SEQ ID NO: 229. | 234 | 97 |
| 722 | Y20298 | Homo sapiens | Human apolipoprotein E mutant protein fragment 11. | 152 | 39 |
| 723 | Y86231 | Homo sapiens | Human secreted protein HLTHR66, SEQ ID NO:146. | 207 | 51 |
| 724 | W75083 | Homo sapiens | Human secreted protein encoded by gene 27 clone HSPAF93. | 685 | 100 |
| 725 | W88627 | Homo sapiens | Secreted protein encoded by gene 94 clone HPMBQ32. | 301 | 73 |
| 726 | Y27868 | Homo sapiens | Human secreted protein encoded by gene No. 107. | 229 | 58 |
| 727 | AK025470 | Homo sapiens | unnamed protein product | 130 | 64 |
| 728 | G02872 | Homo sapiens | Human secreted protein, SEQ ID NO: 6953. | 159 | 46 |
| 729 | Y25776 | Homo sapiens | Human secreted protein encoded from gene 66. | 334 | 43 |
| 730 | AF116661 | Homo sapiens | PRO1438 | 153 | 56 |
| 731 | W48351 | Homo sapiens | Human breast cancer related protein BCRB2. | 106 | 72 |
| 732 | U77589 | Homo sapiens | MHC class II HLA- | 133 | 69 |

TABLE 2

| SEQ ID NO: OF NUCLEOTIDE | ACCESSION NUMBER | SPECIES | DESCRIPTION | SMITH-WATERMAN SCORE | % IDENTITY |
|---|---|---|---|---|---|
| | | | DQ-alpha chain | | |
| 733 | G00357 | Homo sapiens | Human secreted protein, SEQ ID NO: 4438. | 223 | 67 |
| 734 | R28542 | Homo sapiens | Human complement type 1 receptor SCR9. | 152 | 96 |
| 735 | Y27868 | Homo sapiens | Human secreted protein encoded by gene No. 107. | 150 | 65 |
| 736 | AB036706 | Homo sapiens | intelectin | 368 | 76 |
| 737 | Y74042 | Homo sapiens | Human prostate tumor EST fragment derived protein #229. | 206 | 65 |
| 738 | Y36156 | Homo sapiens | Human secreted protein #28. | 153 | 77 |
| 739 | W74802 | Homo sapiens | Human secreted protein encoded by gene 73 clone HSQEL25. | 1751 | 79 |
| 740 | W85614 | Homo sapiens | Secreted protein clone fr473_2. | 224 | 91 |
| 741 | Y13377 | Homo sapiens | Amino acid sequence of protein PRO257. | 394 | 98 |
| 742 | Z69384 | Caenorhabditis elegans | Similarity to Salmonella regulatory protein UHPC (SW:UHPC_SALTY) | 515 | 45 |
| 743 | W47589 | Homo sapiens | T-cell receptor beta-chain. | 681 | 92 |
| 744 | G03786 | Homo sapiens | Human secreted protein, SEQ ID NO: 7867. | 243 | 71 |
| 745 | Y50690 | Homo sapiens | Human Hum4 VL ClaI-HindIII segment encoded protein. | 540 | 81 |
| 746 | U03414 | Rattus norvegicus | neuronal olfactomedin-related ER localized protein | 363 | 67 |
| 747 | G00352 | Homo sapiens | Human secreted protein, SEQ ID NO: 4433. | 84 | 51 |
| 748 | Y02671 | Homo sapiens | Human secreted protein encoded by gene 22 clone HMSJW18. | 145 | 60 |
| 749 | AF026919 | Homo sapiens | amyloid lambda light chain variable region | 557 | 83 |
| 750 | X76732 | Homo sapiens | NEFA protein | 297 | 100 |

# TABLE 2

| SEQ ID NO: OF NUCLEOTIDE | ACCESSION NUMBER | SPECIES | DESCRIPTION | SMITH-WATERMAN SCORE | % IDENTITY |
|---|---|---|---|---|---|
| 751 | R92754 | Homo sapiens | Human growth differentiation factor-12. | 628 | 100 |
| 752 | Y91462 | Homo sapiens | Human secreted protein sequence encoded by gene 12 SEQ ID NO:135. | 597 | 100 |
| 753 | Y66700 | Homo sapiens | Membrane-bound protein PRO1137. | 754 | 99 |
| 754 | G01648 | Homo sapiens | Human secreted protein, SEQ ID NO: 5729. | 281 | 100 |
| 755 | AB040434 | Homo sapiens | hTROY | 752 | 100 |
| 756 | Y28680 | Homo sapiens | Human nm214_3 secreted protein. | 178 | 44 |
| 757 | W75100 | Homo sapiens | Human secreted protein encoded by gene 44 clone HE8CJ26. | 203 | 66 |
| 758 | AF090930 | Homo sapiens | PRO0478 | 87 | 45 |
| 759 | D84336 | Rattus norvegicus | ZOG | 484 | 48 |
| 760 | W88627 | Homo sapiens | Secreted protein encoded by gene 94 clone HPMBQ32. | 150 | 81 |
| 761 | Y48616 | Homo sapiens | Human breast tumour-associated protein 77. | 569 | 70 |
| 762 | Y87320 | Homo sapiens | Human signal peptide containing protein HSPP-97 SEQ ID NO:97. | 918 | 100 |
| 763 | G03655 | Homo sapiens | Human secreted protein, SEQ ID NO: 7736. | 248 | 89 |
| 764 | AF031174 | Homo sapiens | Ig-like membrane protein | 428 | 45 |
| 765 | U08255 | Rattus norvegicus | glutamate receptor delta-1 subunit | 802 | 99 |
| 766 | Y99369 | Homo sapiens | Human PRO1249 (UNQ632) amino acid sequence SEQ ID NO:102. | 4578 | 99 |
| 767 | AK001586 | Homo sapiens | unnamed protein product | 973 | 98 |
| 768 | AC007063 | Arabidopsis thaliana | putative ABC transporter | 126 | 31 |
| 769 | AF303378 | Homo sapiens | sialic acid-specific acetylesterase II | 713 | 100 |
| 770 | G00517 | Homo sapiens | Human secreted protein, SEQ ID NO: 4598. | 90 | 37 |

TABLE 2

| SEQ ID NO: OF NUCLEOTIDE | ACCESSION NUMBER | SPECIES | DESCRIPTION | SMITH-WATERMAN SCORE | % IDENTITY |
|---|---|---|---|---|---|
| 771 | Y59733 | Homo sapiens | Human normal ovarian tissue derived protein 10. | 1253 | 99 |
| 772 | AF132856 | Homo sapiens | suppressor of G2 allele of skp1 homolog | 163 | 86 |
| 773 | AB029482 | Mus musculus | JNK-binding protein JNKBP1 | 1082 | 97 |
| 774 | G02108 | Homo sapiens | Human secreted protein, SEQ ID NO: 6189. | 134 | 62 |
| 775 | AB047818 | Homo sapiens | Soggy | 1239 | 100 |
| 776 | Y66689 | Homo sapiens | Membrane-bound protein PRO1136. | 804 | 99 |
| 777 | Y71107 | Homo sapiens | Human Hydrolase protein-5 (HYDRL-5). | 733 | 99 |
| 778 | AC005626 | Homo sapiens | R29124_1 | 182 | 38 |
| 779 | W88707 | Homo sapiens | Secreted protein encoded by gene 174 clone HE9FB42. | 126 | 56 |
| 780 | G03657 | Homo sapiens | Human secreted protein, SEQ ID NO: 7738. | 455 | 96 |
| 781 | AJ001616 | Mus musculus | myeloid associated differentiation protein | 201 | 36 |
| 782 | Y64942 | Homo sapiens | Human 5' EST related polypeptide SEQ ID NO:1103. | 86 | 65 |
| 783 | AL356276 | Homo sapiens | bA367J7.2.1 (novel Immunoglobulin domains containing protein (isoform 1)) | 845 | 91 |
| 784 | Y00876 | Homo sapiens | Human LAPH-1 protein sequence. | 291 | 43 |
| 785 | G00270 | Homo sapiens | Human secreted protein, SEQ ID NO: 4351. | 603 | 100 |
| 786 | AF154121 | Homo sapiens | sodium-dependent high-affinity dicarboxylate transporter | 864 | 100 |
| 787 | Y29804 | Homo sapiens | Human GABA B receptor subunit HG20 peptide #6. | 83 | 42 |
| 788 | AL080239 | Homo sapiens | bG256O22.1 (similar to IGFALS (insulin-like growth factor binding protein, acid labile subunit)) | 599 | 100 |

# TABLE 2

| SEQ ID NO: OF NUCLEOTIDE | ACCESSION NUMBER | SPECIES | DESCRIPTION | SMITH-WATERMAN SCORE | % IDENTITY |
|---|---|---|---|---|---|
| 789 | AL031856 | Schizosaccharomyces pombe | PUTATIVE GOLGI URIDINE DIPHOSPHATE-N-ACETYLGLUCOSAMINE TRANSPORTER | 192 | 40 |
| 790 | G03448 | Homo sapiens | Human secreted protein, SEQ ID NO: 7529. | 141 | 43 |
| 791 | U81291 | Xenopus laevis | oviductin | 310 | 38 |
| 792 | Y41332 | Homo sapiens | Human secreted protein encoded by gene 25 clone HPIBO48. | 295 | 50 |
| 793 | L20315 | Mus musculus | MPS1 protein | 702 | 77 |
| 794 | G01314 | Homo sapiens | Human secreted protein, SEQ ID NO: 5395. | 91 | 36 |
| 795 | AF003136 | Caenorhabditis elegans | similar to 1-acyl-glycerol-3-phosphate acyltransferases | 122 | 38 |
| 796 | G00637 | Homo sapiens | Human secreted protein, SEQ ID NO: 4718. | 160 | 67 |
| 797 | Y36144 | Homo sapiens | Human secreted protein #16. | 622 | 100 |
| 798 | U09453 | Cricetulus griseus | UDP-N-acetylglucosamine: dolichyl phosphate N-acetylglucosamine 1-phosphate transferase | 178 | 66 |
| 799 | Y76144 | Homo sapiens | Human secreted protein encoded by gene 21. | 633 | 100 |
| 800 | Y73456 | Homo sapiens | Human secreted protein clone yd145_1 protein sequence SEQ ID NO:134. | 413 | 89 |
| 801 | Y86540 | Homo sapiens | Human gene 77-encoded protein fragment, SEQ ID NO:457. | 443 | 96 |
| 802 | U49973 | Homo sapiens | ORF1; MER37; putative transposase similar to pogo element | 311 | 53 |
| 803 | M63573 | Homo sapiens | secreted cyclophilin-like protein | 700 | 88 |
| 804 | AF091622 | Homo sapiens | PHD finger protein | 177 | 100 |

TABLE 2

| SEQ ID NO: OF NUCLEOTIDE | ACCESSION NUMBER | SPECIES | DESCRIPTION | SMITH-WATERMAN SCORE | % IDENTITY |
|---|---|---|---|---|---|
| | | | 3 | | |
| 805 | W37869 | Homo sapiens | Human protein comprising secretory signal amino acid sequence 6. | 381 | 100 |
| 806 | G03556 | Homo sapiens | Human secreted protein, SEQ ID NO: 7637. | 221 | 72 |
| 807 | AF178941 | Homo sapiens | ATP-binding cassette sub-family A member 2 | 583 | 87 |
| 808 | Y91385 | Homo sapiens | Human secreted protein sequence encoded by gene 40 SEQ ID NO:106. | 786 | 100 |
| 809 | Y00826 | Rattus norvegicus | gp210 (AA 1-1886) | 169 | 83 |
| 810 | G03143 | Homo sapiens | Human secreted protein, SEQ ID NO: 7224. | 328 | 100 |
| 811 | W00870 | Homo sapiens | Polycystic kidney disease 1 (PKD1) polypeptide. | 22446 | 99 |
| 812 | Y73434 | Homo sapiens | Human secreted protein clone yd51_1 protein sequence SEQ ID NO:90. | 417 | 90 |
| 813 | AB031996 | Ralstonia sp. KN1 | ferredoxin-like protein | 94 | 44 |
| 814 | AF201734 | Mus musculus | testis specific serine kinase-3 | 800 | 87 |
| 815 | Y01181 | Homo sapiens | Polypeptide fragment encoded by gene 12. | 68 | 55 |
| 816 | Y76166 | Homo sapiens | Human secreted protein encoded by gene 43. | 724 | 94 |
| 817 | AL109827 | Homo sapiens | dJ309K20.2 (acrosomal protein ACR55 (similar to rat sperm antigen 4 (SPAG4))) | 639 | 84 |
| 818 | M62829 | Homo sapiens | ETR103 | 137 | 53 |
| 819 | Y38422 | Homo sapiens | Human secreted protein. | 526 | 100 |
| 820 | AF119815 | Homo sapiens | G-protein-coupled receptor | 561 | 79 |
| 821 | Y87101 | Homo sapiens | Human secreted protein sequence SEQ ID NO:140. | 628 | 100 |
| 822 | M91463 | Homo sapiens | glucose transporter | 213 | 79 |

# TABLE 2

| SEQ ID NO: OF NUCLEOTIDE | ACCESSION NUMBER | SPECIES | DESCRIPTION | SMITH-WATERMAN SCORE | % IDENTITY |
|---|---|---|---|---|---|
| 823 | L34938 | Rattus norvegicus | ionotropic glutamate receptor | 618 | 90 |
| 824 | W17846 | Homo sapiens | Cytosolic phospholipase A2/B (clone 19b product). | 209 | 64 |
| 825 | Y66722 | Homo sapiens | Membrane-bound protein PRO1104. | 221 | 67 |
| 826 | G02493 | Homo sapiens | Human secreted protein, SEQ ID NO: 6574. | 138 | 72 |
| 827 | Y91423 | Homo sapiens | Human secreted protein sequence encoded by gene 11 SEQ ID NO:144. | 671 | 54 |
| 828 | U78090 | Rattus norvegicus | potassium channel regulator 1 | 502 | 80 |
| 829 | U08813 | Oryctolagus cuniculus | 597 aa protein related to Na/glucose cotransporters | 906 | 84 |
| 830 | AJ272063 | Homo sapiens | vanilloid receptor 1 | 630 | 90 |
| 831 | U36898 | Rattus norvegicus | pheromone receptor VN6 | 135 | 52 |
| 832 | Z46973 | Homo sapiens | phosphatidylinositol 3-kinase | 396 | 80 |
| 833 | Y95433 | Homo sapiens | Human calcium channel SOC-2/CRAC-1 C-terminal polypeptide. | 747 | 99 |
| 834 | AF132856 | Homo sapiens | suppressor of G2 allele of skp1 homolog | 163 | 86 |
| 835 | AC006042 | Homo sapiens | supported by human ESTs AI681256.1(NID:g4891438),N32168.1(NID:g1152567), and genscan | 195 | 87 |
| 836 | B01247 | Homo sapiens | Human HE6 receptor. | 371 | 45 |
| 837 | G03788 | Homo sapiens | Human secreted protein, SEQ ID NO: 7869. | 196 | 59 |
| 838 | U70136 | Homo sapiens | megakaryocyte stimulating factor; MSF | 6954 | 98 |
| 839 | AF017153 | Mus musculus | putative RNA helicase and RNA dependent ATPase | 178 | 51 |
| 840 | Y31830 | Homo sapiens | Human adult brain secreted protein nh899_8. | 244 | 56 |

# TABLE 2

| SEQ ID NO: OF NUCLEOTIDE | ACCESSION NUMBER | SPECIES | DESCRIPTION | SMITH-WATERMAN SCORE | % IDENTITY |
|---|---|---|---|---|---|
| 841 | Y27593 | Homo sapiens | Human secreted protein encoded by gene No. 27. | 437 | 81 |
| 842 | G01984 | Homo sapiens | Human secreted protein, SEQ ID NO: 6065. | 196 | 74 |
| 843 | AL008723 | Homo sapiens | dJ90G24.4 (SAAT1 (low affinity sodium glucose cotransporter (sodium:solute symporter family))) | 183 | 92 |
| 844 | AF068065 | Cryptosporidium parvum | GP900; mucin-like glycoprotein | 263 | 47 |
| 845 | Y00815 | Homo sapiens | put. LAR preprotein (AA -16 to 1881) | 341 | 100 |
| 846 | Y06816 | Homo sapiens | Human Notch2 (humN2) protein sequence. | 1224 | 99 |
| 847 | AF104923 | Homo sapiens | putative transcription factor | 293 | 95 |
| 848 | Y09945 | Rattus norvegicus | putative integral membrane transport protein | 589 | 53 |
| 849 | AL157874 | Schizosaccharomyces pombe | similar to yeast SCT1 suppressor of a choline transport mutant | 146 | 40 |
| 850 | R71003 | Homo sapiens | Human neuronal calcium channel subunit alpha 1c-1. | 141 | 89 |
| 851 | X75756 | Homo sapiens | protein kinase C mu | 318 | 90 |
| 852 | AF142676 | Drosophila melanogaster | sodium-hydrogen exchanger NHE1 | 366 | 48 |
| 853 | Y45381 | Homo sapiens | Human secreted protein fragment encoded from gene 28. | 139 | 73 |
| 854 | G03789 | Homo sapiens | Human secreted protein, SEQ ID NO: 7870. | 121 | 60 |
| 855 | U65409 | Yarrowia lipolytica | Sla2p | 109 | 25 |
| 856 | M19419 | Mus musculus | proline-rich salivary protein | 109 | 36 |
| 857 | Y99355 | Homo sapiens | Human PRO1295 (UNQ664) amino acid sequence SEQ ID NO:54. | 667 | 98 |
| 858 | W19919 | Homo sapiens | Human Ksr-1 (kinase suppressor of Ras). | 211 | 86 |
| 859 | Y95436 | Homo sapiens | Human calcium | 764 | 84 |

# TABLE 2

| SEQ ID NO: OF NUCLEOTIDE | ACCESSION NUMBER | SPECIES | DESCRIPTION | SMITH-WATERMAN SCORE | % IDENTITY |
|---|---|---|---|---|---|
| | | | channel SOC-3/CRAC-2. | | |
| 860 | AF070066 | Mus musculus | Citron-K kinase | 628 | 97 |
| 861 | AF286095 | Homo sapiens | IL-22 receptor | 933 | 100 |
| 862 | AF020195 | Mus musculus | pancreas sodium bicarbonate cotransporter | 475 | 68 |
| 863 | G03712 | Homo sapiens | Human secreted protein, SEQ ID NO: 7793. | 240 | 100 |
| 864 | AF195092 | Homo sapiens | sialic acid-binding immunoglobulin-like lectin-8 | 288 | 87 |
| 865 | AF208110 | Homo sapiens | IL-17 receptor homolog precursor | 2688 | 99 |
| 866 | L42338 | Mus musculus | sodium channel 25 | 733 | 98 |
| 867 | G02360 | Homo sapiens | Human secreted protein, SEQ ID NO: 6441. | 101 | 70 |
| 868 | AF065215 | Homo sapiens | cytosolic phospholipase A2 beta | 290 | 42 |
| 869 | L43631 | Homo sapiens | scaffold attachment factor B | 106 | 95 |
| 870 | G03034 | Homo sapiens | Human secreted protein, SEQ ID NO: 7115. | 108 | 54 |
| 871 | Z21514 | Rattus norvegicus | integral membrane glycoprotein | 84 | 47 |
| 872 | AF097518 | Homo sapiens | liver-specific transporter | 147 | 40 |
| 873 | AF288223 | Drosophila melanogaster | Crossveinless 2 | 136 | 39 |
| 874 | U90126 | Bos taurus | ABC transporter | 245 | 36 |
| 875 | AF099988 | Mus musculus | Ste-20 related kinase SPAK | 103 | 34 |
| 876 | Y70400 | Homo sapiens | Human cell-signalling protein-2. | 220 | 86 |
| 877 | Y36300 | Homo sapiens | Human secreted protein encoded by gene 77. | 1863 | 99 |
| 878 | AF151074 | Homo sapiens | HSPC240 | 193 | 29 |
| 879 | Y94951 | Homo sapiens | Human secreted protein clone dw78_1 protein sequence SEQ ID NO:108. | 251 | 89 |
| 880 | AF165310 | Homo sapiens | ATP cassette binding transporter 1 | 231 | 31 |
| 881 | AF252281 | Mus musculus | Kelch-like 1 protein | 256 | 58 |

# TABLE 2

| SEQ ID NO: OF NUCLEOTIDE | ACCESSION NUMBER | SPECIES | DESCRIPTION | SMITH-WATERMAN SCORE | % IDENTITY |
|---|---|---|---|---|---|
| 882 | Y00931 | Homo sapiens | Prostate-tumour derived antigen #4. | 1039 | 98 |
| 883 | Y27576 | Homo sapiens | Human secreted protein encoded by gene No. 10. | 394 | 96 |
| 884 | U00009 | Escherichia coli | yeeF | 153 | 30 |
| 885 | Y57945 | Homo sapiens | Human transmembrane protein HTMPN-69. | 1543 | 100 |
| 886 | Y28678 | Homo sapiens | Human cw272_7 secreted protein. | 375 | 60 |
| 887 | W95349 | Homo sapiens | Human foetal brain secreted protein fh170_7. | 377 | 89 |
| 888 | Y87329 | Homo sapiens | Human signal peptide containing protein HSPP-106 SEQ ID NO:106. | 285 | 89 |
| 889 | AL121845 | Homo sapiens | dJ583P15.5.1 (novel protein (isoform 1)) | 1399 | 99 |
| 890 | R75181 | Homo sapiens | Partial peptide of human HMW kininogen fragment 1.2. | 100 | 29 |
| 891 | AF105365 | Homo sapiens | K-Cl cotransporter KCC4 | 680 | 100 |
| 892 | Y91644 | Homo sapiens | Human secreted protein sequence encoded by gene 43 SEQ ID NO:317. | 673 | 95 |
| 893 | S52051 | Rattus sp. | neurotransmitter transporter | 656 | 99 |
| 894 | S52051 | Rattus sp. | neurotransmitter transporter | 617 | 94 |
| 895 | R47120 | Homo sapiens | Partial human H13 polypeptide. | 343 | 60 |
| 896 | Z98046 | Homo sapiens | dJ1409.2 (Melanoma-Associated Antigen MAGE LIKE) | 332 | 49 |
| 897 | AJ006203 | Oryctolagus cuniculus | capacitative calcium entry channel 2 | 740 | 99 |
| 898 | AF156547 | Mus musculus | putative E1-E2 ATPase | 769 | 95 |
| 899 | AC004076 | Homo sapiens | R30217_1 | 788 | 98 |
| 900 | D00099 | Homo sapiens | Na,K-ATPase alpha-subunit | 753 | 94 |
| 901 | R27648 | Homo sapiens | Human calcium channel 27980/10. | 536 | 85 |
| 902 | Y57955 | Homo sapiens | Human transmembrane protein HTMPN-79. | 606 | 100 |
| 903 | AF155913 | Mus musculus | putative E1-E2 ATPase | 1039 | 85 |

# TABLE 2

| SEQ ID NO: OF NUCLEOTIDE | ACCESSION NUMBER | SPECIES | DESCRIPTION | SMITH-WATERMAN SCORE | % IDENTITY |
|---|---|---|---|---|---|
| 904 | Y73446 | Homo sapiens | Human secreted protein clone yc27_1 protein sequence SEQ ID NO:114. | 369 | 66 |
| 905 | Y94903 | Homo sapiens | Human secreted protein clone pt332_1 protein sequence SEQ ID NO:12. | 3777 | 100 |
| 906 | AB032470 | Homo sapiens | seven transmembrane protein TM7SF3 | 2124 | 100 |
| 907 | G00517 | Homo sapiens | Human secreted protein, SEQ ID NO: 4598. | 90 | 50 |
| 908 | AF010144 | Homo sapiens | neuronal thread protein AD7c-NTP | 270 | 65 |
| 909 | AF263912 | Streptomyces noursei | NysA | 113 | 25 |
| 910 | Y53051 | Homo sapiens | Human secreted protein clone dd119_4 protein sequence SEQ ID NO:108. | 843 | 49 |
| 911 | Y76179 | Homo sapiens | Human secreted protein encoded by gene 56. | 634 | 100 |
| 912 | G00352 | Homo sapiens | Human secreted protein, SEQ ID NO: 4433. | 229 | 71 |
| 913 | U93569 | Homo sapiens | p40 | 110 | 32 |
| 914 | G02639 | Homo sapiens | Human secreted protein, SEQ ID NO: 6720. | 65 | 46 |
| 915 | Y94951 | Homo sapiens | Human secreted protein clone dw78_1 protein sequence SEQ ID NO:108. | 100 | 38 |
| 916 | G03263 | Homo sapiens | Human secreted protein, SEQ ID NO: 7344. | 80 | 47 |
| 917 | W74887 | Homo sapiens | Human secreted protein encoded by gene 160 clone HCELB21. | 273 | 69 |
| 918 | Y73464 | Homo sapiens | Human secreted protein clone yl4_1 protein sequence SEQ ID NO:150. | 982 | 90 |
| 919 | AF064801 | Homo sapiens | multiple membrane spanning receptor TRC8 | 551 | 32 |

# TABLE 2

| SEQ ID NO: OF NUCLEOTIDE | ACCESSION NUMBER | SPECIES | DESCRIPTION | SMITH-WATERMAN SCORE | % IDENTITY |
|---|---|---|---|---|---|
| 920 | Y87335 | Homo sapiens | Human signal peptide containing protein HSPP-112 SEQ ID NO:112. | 622 | 99 |
| 921 | AK000496 | Homo sapiens | unnamed protein product | 342 | 74 |
| 922 | Y41360 | Homo sapiens | Human secreted protein encoded by gene 53 clone HJPAD75. | 367 | 100 |
| 923 | G02872 | Homo sapiens | Human secreted protein, SEQ ID NO: 6953. | 328 | 75 |
| 924 | Y53881 | Homo sapiens | A suppressor of cytokine signalling protein designated HSCOP-1. | 1489 | 100 |
| 925 | AC004144 | Homo sapiens | R34001_1 | 193 | 60 |
| 926 | AF119851 | Homo sapiens | PRO1722 | 153 | 82 |
| 927 | G02654 | Homo sapiens | Human secreted protein, SEQ ID NO: 6735. | 82 | 57 |
| 928 | Y30819 | Homo sapiens | Human secreted protein encoded from gene 9. | 264 | 33 |
| 929 | G01691 | Homo sapiens | Human secreted protein, SEQ ID NO: 5772. | 66 | 43 |
| 930 | AF187845 | Homo sapiens | small protein effector 1 of Cdc42 | 431 | 100 |
| 931 | AL390114 | Leishmania major | extremely cysteine/valine rich protein | 113 | 40 |
| 932 | AL080239 | Homo sapiens | bG256022.1 (similar to IGFALS (insulin-like growth factor binding protein, acid labile subunit)) | 1451 | 97 |
| 933 | W85613 | Homo sapiens | Secreted protein clone fm60_1. | 234 | 100 |
| 934 | AF009243 | Homo sapiens | proline-rich Gla protein 2 | 223 | 42 |
| 935 | G03789 | Homo sapiens | Human secreted protein, SEQ ID NO: 7870. | 271 | 66 |
| 936 | AK000385 | Homo sapiens | unnamed protein product | 193 | 64 |
| 937 | AF010144 | Homo sapiens | neuronal thread protein AD7c-NTP | 270 | 65 |
| 938 | AF119851 | Homo sapiens | PRO1722 | 170 | 71 |
| 939 | Y07922 | Homo sapiens | Human secreted protein fragment | 226 | 95 |

# TABLE 2

| SEQ ID NO: OF NUCLEOTIDE | ACCESSION NUMBER | SPECIES | DESCRIPTION | SMITH-WATERMAN SCORE | % IDENTITY |
|---|---|---|---|---|---|
| | | | encoded from gene 71. | | |
| 940 | Y41712 | Homo sapiens | Human PRO724 protein sequence. | 653 | 96 |
| 941 | AF010144 | Homo sapiens | neuronal thread protein AD7c-NTP | 310 | 64 |
| 942 | Y45318 | Homo sapiens | Human secreted protein fragment encoded from gene 18. | 502 | 98 |
| 943 | Y07899 | Homo sapiens | Human secreted protein fragment encoded from gene 48. | 309 | 98 |
| 944 | X92485 | Plasmodium vivax | pval | 185 | 51 |
| 945 | AJ289133 | Mus musculus | chondroitin 4-O-sulfotransferase | 565 | 43 |
| 946 | AF151074 | Homo sapiens | HSPC240 | 1337 | 99 |
| 947 | U40829 | Saccharomyces cerevisiae | Weak similarity near C-terminus to RNA Polymerase beta subunit (Swiss Prot. accession number P11213) and CCAAT-binding transcription factor (PIR accession number A36368) | 361 | 50 |
| 948 | Y87285 | Homo sapiens | Human signal peptide containing protein HSPP-62 SEQ ID NO:62. | 348 | 82 |
| 949 | Y86230 | Homo sapiens | Human secreted protein HKFBC53, SEQ ID NO:145. | 368 | 80 |
| 950 | AJ010346 | Homo sapiens | RING-H2 | 333 | 87 |
| 951 | Z56281 | Homo sapiens | interferon regulatory factor 3 | 1573 | 81 |
| 952 | Y57896 | Homo sapiens | Human transmembrane protein HTMPN-20. | 421 | 100 |
| 953 | G03789 | Homo sapiens | Human secreted protein, SEQ ID NO: 7870. | 135 | 55 |
| 954 | Y87103 | Homo sapiens | Human secreted protein sequence SEQ ID NO:142. | 83 | 50 |
| 955 | Y87345 | Homo sapiens | Human signal peptide containing protein HSPP-122 SEQ ID NO:122. | 885 | 99 |
| 956 | X81479 | Homo sapiens | EMR1 | 1148 | 99 |

# TABLE 2

| SEQ ID NO: OF NUCLEOTIDE | ACCESSION NUMBER | SPECIES | DESCRIPTION | SMITH-WATERMAN SCORE | % IDENTITY |
|---|---|---|---|---|---|
| 957 | AF175406 | Homo sapiens | transient receptor potential 4 | 4061 | 99 |
| 958 | G03789 | Homo sapiens | Human secreted protein, SEQ ID NO: 7870. | 276 | 73 |
| 959 | M63274 | Plasmodium falciparum | malaria antigen | 77 | 38 |
| 960 | Y78795 | Homo sapiens | Human antizuai-2 (AZ-2) amino acid sequence. | 3384 | 83 |
| 961 | AL133469 | Streptomyces coelicolor A3(2) | putative secreted proline-rich protein | 139 | 41 |
| 962 | G03787 | Homo sapiens | Human secreted protein, SEQ ID NO: 7868. | 232 | 72 |
| 963 | W74828 | Homo sapiens | Human secreted protein encoded by gene 100 clone HLQAB52. | 1016 | 99 |
| 964 | W48351 | Homo sapiens | Human breast cancer related protein BCRB2. | 226 | 58 |
| 965 | X63893 | Sus scrofa | alpha-stimulatory subunit of GTP-binding protein | 319 | 86 |
| 966 | AB033019 | Homo sapiens | KIAA1193 protein | 245 | 97 |
| 967 | Y36156 | Homo sapiens | Human secreted protein #28. | 223 | 85 |
| 968 | AF119851 | Homo sapiens | PRO1722 | 188 | 69 |
| 969 | Y15224 | Homo sapiens | Human receptor protein (HURP) 3 amino acid sequence. | 214 | 42 |
| 970 | G02754 | Homo sapiens | Human secreted protein, SEQ ID NO: 6835. | 81 | 62 |
| 971 | U22376 | Homo sapiens | alternatively spliced product using exon 13A | 212 | 81 |
| 972 | W74870 | Homo sapiens | Human secreted protein encoded by gene 142 clone HTWCB92. | 164 | 81 |
| 973 | Y30817 | Homo sapiens | Human secreted protein encoded from gene 7. | 717 | 98 |
| 974 | AF079529 | Homo sapiens | cAMP-specific phosphodiesterase 8B; PDE8B1; 3',5'-cyclic nucleotide phosphodiesterase | 2353 | 96 |
| 975 | AF099028 | Drosophila | putative | 1061 | 52 |

TABLE 2

| SEQ ID NO: OF NUCLEOTIDE | ACCESSION NUMBER | SPECIES | DESCRIPTION | SMITH-WATERMAN SCORE | % IDENTITY |
|---|---|---|---|---|---|
| | | melanogaster | transmembrane protein cmp44E | | |
| 976 | G03786 | Homo sapiens | Human secreted protein, SEQ ID NO: 7867. | 179 | 72 |
| 977 | Y22495 | Homo sapiens | Human secreted protein sequence clone ch4_11. | 1629 | 100 |
| 978 | W74813 | Homo sapiens | Human secreted protein encoded by gene 85 clone HSDFV29. | 722 | 92 |
| 979 | AK023408 | Homo sapiens | unnamed protein product | 974 | 96 |
| 980 | AF229178 | Homo sapiens | leucine rich repeat and death domain containing protein | 276 | 67 |
| 981 | G03797 | Homo sapiens | Human secreted protein, SEQ ID NO: 7878. | 198 | 56 |
| 982 | W74831 | Homo sapiens | Human secreted protein encoded by gene 103 clone HEBDJ82. | 153 | 100 |
| 983 | G01335 | Homo sapiens | Human secreted protein, SEQ ID NO: 5416. | 157 | 96 |
| 984 | Y73436 | Homo sapiens | Human secreted protein clone ye43_1 protein sequence SEQ ID NO:94. | 450 | 100 |
| 985 | G00354 | Homo sapiens | Human secreted protein, SEQ ID NO: 4435. | 96 | 58 |
| 986 | Y41712 | Homo sapiens | Human PRO724 protein sequence. | 639 | 88 |
| 987 | Y57896 | Homo sapiens | Human transmembrane protein HTMPN-20. | 421 | 100 |
| 988 | Y66691 | Homo sapiens | Membrane-bound protein PRO809. | 716 | 65 |
| 989 | AF090943 | Homo sapiens | PRO0659 | 926 | 100 |
| 990 | G00403 | Homo sapiens | Human secreted protein, SEQ ID NO: 4484. | 80 | 46 |
| 991 | G03411 | Homo sapiens | Human secreted protein, SEQ ID NO: 7492. | 62 | 57 |
| 992 | G00270 | Homo sapiens | Human secreted protein, SEQ ID NO: 4351. | 143 | 96 |
| 993 | AF026246 | Homo sapiens | HERV-E integrase | 361 | 80 |
| 994 | Y36421 | Homo sapiens | Fragment of human | 83 | 37 |

# TABLE 2

| SEQ ID NO: OF NUCLEOTIDE | ACCESSION NUMBER | SPECIES | DESCRIPTION | SMITH-WATERMAN SCORE | % IDENTITY |
|---|---|---|---|---|---|
| | | | secreted protein encoded by gene 8. | | |
| 995 | U22376 | Homo sapiens | alternatively spliced product using exon 13A | 175 | 78 |
| 996 | G03790 | Homo sapiens | Human secreted protein, SEQ ID NO: 7871. | 87 | 35 |
| 997 | G00397 | Homo sapiens | Human secreted protein, SEQ ID NO: 4478. | 149 | 61 |
| 998 | J02642 | Homo sapiens | glyceraldehyde 3-phosphate dehydrogenase (EC 1.2.1.12) | 429 | 69 |
| 999 | AF119851 | Homo sapiens | PRO1722 | 204 | 50 |
| 1000 | Y91423 | Homo sapiens | Human secreted protein sequence encoded by gene 11 SEQ ID NO:144. | 393 | 53 |
| 1001 | Y66695 | Homo sapiens | Membrane-bound protein PRO1344. | 1183 | 87 |
| 1002 | AF090931 | Homo sapiens | PRO0483 | 149 | 68 |
| 1003 | Y33261 | Homo sapiens | Human p99 protein. | 314 | 59 |
| 1004 | U11494 | Mus musculus | protein kinase | 360 | 77 |
| 1005 | AK021848 | Homo sapiens | unnamed protein product | 186 | 69 |
| 1006 | Y13892 | Homo sapiens | PI-3 kinase | 233 | 97 |
| 1007 | W48351 | Homo sapiens | Human breast cancer related protein BCRB2. | 144 | 65 |
| 1008 | G03793 | Homo sapiens | Human secreted protein, SEQ ID NO: 7874. | 202 | 67 |
| 1009 | U91682 | Aedes aegypti | vitelline membrane protein homolog | 88 | 42 |

# TABLE 3

| SEQ ID NO: OF NUCLEOTIDE | SEQ ID NO: OF AMINO ACID | SEQ ID NO: IN USSN 09/491,404 | START NUCLEOTIDE OF CODING REGION | STOP NUCLEOTIDE OF CODING REGION |
|---|---|---|---|---|
| 1 | 1010 | 100 | 299 | 535 |
| 2 | 1011 | 1002 | 19 | 267 |
| 3 | 1012 | 1003 | 31 | 423 |
| 4 | 1013 | 1007 | 148 | 840 |
| 5 | 1014 | 1009 | 139 | 318 |
| 6 | 1015 | 1010 | 413 | 748 |
| 7 | 1016 | 1012 | 357 | 154 |
| 8 | 1017 | 1014 | 133 | 285 |
| 9 | 1018 | 1016 | 61 | 441 |
| 10 | 1019 | 102 | 269 | 832 |
| 11 | 1020 | 1021 | 148 | 342 |
| 12 | 1021 | 1022 | 45 | 452 |
| 13 | 1022 | 1035 | 222 | 779 |
| 14 | 1023 | 1038 | 222 | 779 |
| 15 | 1024 | 1042 | 735 | 517 |
| 16 | 1025 | 1049 | 120 | 320 |
| 17 | 1026 | 1055 | 195 | 395 |
| 18 | 1027 | 1061 | 13 | 189 |
| 19 | 1028 | 1070 | 972 | 1109 |
| 20 | 1029 | 1071 | 1504 | 1686 |
| 21 | 1030 | 1077 | 425 | 574 |
| 22 | 1031 | 108 | 46 | 501 |
| 23 | 1032 | 1088 | 1949 | 7240 |
| 24 | 1033 | 1092 | 119 | 571 |
| 25 | 1034 | 1095 | 118 | 564 |
| 26 | 1035 | 1096 | 110 | 373 |
| 27 | 1036 | 1098 | 66 | 353 |
| 28 | 1037 | 1099 | 1 | 417 |
| 29 | 1038 | 11 | 764 | 573 |
| 30 | 1039 | 1100 | 157 | 1014 |
| 31 | 1040 | 1102 | 1526 | 1813 |
| 32 | 1041 | 1103 | 1529 | 1338 |
| 33 | 1042 | 1104 | 685 | 1929 |
| 34 | 1043 | 1105 | 887 | 744 |
| 35 | 1044 | 1110 | 880 | 443 |
| 36 | 1045 | 1111 | 696 | 538 |
| 37 | 1046 | 1113 | 52 | 1272 |
| 38 | 1047 | 1117 | 1357 | 554 |
| 39 | 1048 | 1118 | 1478 | 1654 |
| 40 | 1049 | 112 | 482 | 712 |
| 41 | 1050 | 1121 | 3 | 1424 |
| 42 | 1051 | 1130 | 131 | 271 |
| 43 | 1052 | 1132 | 849 | 151 |
| 44 | 1053 | 1137 | 265 | 705 |
| 45 | 1054 | 1138 | 13 | 381 |
| 46 | 1055 | 1140 | 51 | 416 |
| 47 | 1056 | 1146 | 2389 | 2541 |
| 48 | 1057 | 1148 | 1517 | 738 |
| 49 | 1058 | 115 | 179 | 334 |
| 50 | 1059 | 1154 | 68 | 358 |

# TABLE 3

| SEQ ID NO: OF NUCLEOTIDE | SEQ ID NO: OF AMINO ACID | SEQ ID NO: IN USSN 09/491,404 | START NUCLEOTIDE OF CODING REGION | STOP NUCLEOTIDE OF CODING REGION |
|---|---|---|---|---|
| 51 | 1060 | 1155 | 34 | 330 |
| 52 | 1061 | 1157 | 242 | 433 |
| 53 | 1062 | 1160 | 410 | 856 |
| 54 | 1063 | 1161 | 154 | 342 |
| 55 | 1064 | 1163 | 202 | 477 |
| 56 | 1065 | 1167 | 72 | 272 |
| 57 | 1066 | 117 | 235 | 2 |
| 58 | 1067 | 1170 | 47 | 211 |
| 59 | 1068 | 1176 | 16 | 159 |
| 60 | 1069 | 1177 | 135 | 326 |
| 61 | 1070 | 118 | 1248 | 1466 |
| 62 | 1071 | 1183 | 431 | 886 |
| 63 | 1072 | 1187 | 191 | 529 |
| 64 | 1073 | 1189 | 1303 | 1148 |
| 65 | 1074 | 119 | 380 | 613 |
| 66 | 1075 | 1190 | 514 | 1272 |
| 67 | 1076 | 1192 | 1529 | 1338 |
| 68 | 1077 | 1197 | 93 | 533 |
| 69 | 1078 | 1199 | 227 | 391 |
| 70 | 1079 | 1202 | 117 | 407 |
| 71 | 1080 | 1204 | 12 | 413 |
| 72 | 1081 | 1205 | 49 | 603 |
| 73 | 1082 | 1216 | 487 | 1341 |
| 74 | 1083 | 1217 | 982 | 764 |
| 75 | 1084 | 1228 | 99 | 266 |
| 76 | 1085 | 1230 | 973 | 770 |
| 77 | 1086 | 1233 | 233 | 418 |
| 78 | 1087 | 1234 | 2959 | 2078 |
| 79 | 1088 | 1235 | 112 | 1542 |
| 80 | 1089 | 1239 | 3019 | 2822 |
| 81 | 1090 | 1242 | 1335 | 781 |
| 82 | 1091 | 1248 | 29 | 169 |
| 83 | 1092 | 125 | 542 | 405 |
| 84 | 1093 | 1250 | 1381 | 1572 |
| 85 | 1094 | 1252 | 480 | 226 |
| 86 | 1095 | 1255 | 19 | 285 |
| 87 | 1096 | 1259 | 165 | 638 |
| 88 | 1097 | 126 | 627 | 364 |
| 89 | 1098 | 1260 | 289 | 462 |
| 90 | 1099 | 1262 | 138 | 353 |
| 91 | 1100 | 1264 | 1159 | 1299 |
| 92 | 1101 | 1266 | 13 | 402 |
| 93 | 1102 | 1269 | 296 | 805 |
| 94 | 1103 | 127 | 212 | 397 |
| 95 | 1104 | 1270 | 126 | 374 |
| 96 | 1105 | 1272 | 2025 | 2396 |
| 97 | 1106 | 1273 | 1367 | 624 |
| 98 | 1107 | 1274 | 1108 | 746 |
| 99 | 1108 | 1275 | 919 | 1077 |
| 100 | 1109 | 1279 | 496 | 1272 |

# TABLE 3

| SEQ ID NO: OF NUCLEOTIDE | SEQ ID NO: OF AMINO ACID | SEQ ID NO: IN USSN 09/491,404 | START NUCLEOTIDE OF CODING REGION | STOP NUCLEOTIDE OF CODING REGION |
|---|---|---|---|---|
| 101 | 1110 | 1283 | 265 | 125 |
| 102 | 1111 | 1287 | 107 | 385 |
| 103 | 1112 | 1297 | 333 | 545 |
| 104 | 1113 | 13 | 187 | 47 |
| 105 | 1114 | 130 | 126 | 290 |
| 106 | 1115 | 1306 | 323 | 75 |
| 107 | 1116 | 1308 | 457 | 891 |
| 108 | 1117 | 1311 | 258 | 674 |
| 109 | 1118 | 1315 | 242 | 823 |
| 110 | 1119 | 1317 | 82 | 435 |
| 111 | 1120 | 1319 | 781 | 3306 |
| 112 | 1121 | 1323 | 1402 | 1671 |
| 113 | 1122 | 1329 | 279 | 665 |
| 114 | 1123 | 1336 | 37 | 765 |
| 115 | 1124 | 1337 | 177 | 389 |
| 116 | 1125 | 1338 | 887 | 744 |
| 117 | 1126 | 1339 | 248 | 724 |
| 118 | 1127 | 1341 | 298 | 525 |
| 119 | 1128 | 1342 | 26 | 445 |
| 120 | 1129 | 1344 | 23 | 370 |
| 121 | 1130 | 1345 | 160 | 402 |
| 122 | 1131 | 1351 | 2737 | 2600 |
| 123 | 1132 | 1353 | 655 | 792 |
| 124 | 1133 | 1354 | 94 | 354 |
| 125 | 1134 | 1356 | 679 | 849 |
| 126 | 1135 | 1358 | 679 | 849 |
| 127 | 1136 | 1359 | 32 | 346 |
| 128 | 1137 | 1361 | 271 | 426 |
| 129 | 1138 | 1362 | 637 | 1197 |
| 130 | 1139 | 1363 | 24 | 350 |
| 131 | 1140 | 1364 | 119 | 367 |
| 132 | 1141 | 1368 | 111 | 284 |
| 133 | 1142 | 1377 | 1221 | 1358 |
| 134 | 1143 | 1378 | 643 | 470 |
| 135 | 1144 | 138 | 99 | 539 |
| 136 | 1145 | 1382 | 994 | 686 |
| 137 | 1146 | 1384 | 34 | 264 |
| 138 | 1147 | 1386 | 124 | 477 |
| 139 | 1148 | 1389 | 1197 | 1 |
| 140 | 1149 | 139 | 94 | 294 |
| 141 | 1150 | 1390 | 1262 | 1053 |
| 142 | 1151 | 1393 | 1182 | 1325 |
| 143 | 1152 | 1394 | 1351 | 1542 |
| 144 | 1153 | 1395 | 229 | 411 |
| 145 | 1154 | 1396 | 923 | 1147 |
| 146 | 1155 | 1397 | 49 | 252 |
| 147 | 1156 | 1398 | 684 | 863 |
| 148 | 1157 | 1399 | 2613 | 286 |
| 149 | 1158 | 14 | 997 | 758 |
| 150 | 1159 | 1403 | 396 | 1 |

# TABLE 3

| SEQ ID NO: OF NUCLEOTIDE | SEQ ID NO: OF AMINO ACID | SEQ ID NO: IN USSN 09/491,404 | START NUCLEOTIDE OF CODING REGION | STOP NUCLEOTIDE OF CODING REGION |
|---|---|---|---|---|
| 151 | 1160 | 1406 | 735 | 1235 |
| 152 | 1161 | 1407 | 967 | 716 |
| 153 | 1162 | 1408 | 75 | 314 |
| 154 | 1163 | 1409 | 101 | 313 |
| 155 | 1164 | 141 | 384 | 551 |
| 156 | 1165 | 1414 | 242 | 532 |
| 157 | 1166 | 142 | 158 | 15 |
| 158 | 1167 | 1421 | 604 | 1425 |
| 159 | 1168 | 1422 | 1146 | 1835 |
| 160 | 1169 | 1423 | 2657 | 3295 |
| 161 | 1170 | 1424 | 315 | 163 |
| 162 | 1171 | 1426 | 39 | 509 |
| 163 | 1172 | 1427 | 892 | 686 |
| 164 | 1173 | 1428 | 395 | 619 |
| 165 | 1174 | 1430 | 284 | 514 |
| 166 | 1175 | 1432 | 178 | 2 |
| 167 | 1176 | 1433 | 1136 | 972 |
| 168 | 1177 | 1435 | 1283 | 1540 |
| 169 | 1178 | 1436 | 1669 | 2235 |
| 170 | 1179 | 144 | 55 | 219 |
| 171 | 1180 | 1440 | 363 | 121 |
| 172 | 1181 | 1441 | 1991 | 2197 |
| 173 | 1182 | 1443 | 1765 | 3054 |
| 174 | 1183 | 1445 | 1023 | 865 |
| 175 | 1184 | 1446 | 5692 | 5859 |
| 176 | 1185 | 1447 | 2959 | 2078 |
| 177 | 1186 | 1448 | 775 | 945 |
| 178 | 1187 | 1451 | 858 | 1430 |
| 179 | 1188 | 1453 | 1370 | 723 |
| 180 | 1189 | 1455 | 480 | 1007 |
| 181 | 1190 | 1457 | 278 | 451 |
| 182 | 1191 | 1459 | 824 | 561 |
| 183 | 1192 | 1460 | 56 | 463 |
| 184 | 1193 | 1461 | 184 | 480 |
| 185 | 1194 | 1462 | 486 | 635 |
| 186 | 1195 | 1465 | 319 | 492 |
| 187 | 1196 | 1466 | 398 | 3 |
| 188 | 1197 | 1468 | 262 | 453 |
| 189 | 1198 | 1476 | 526 | 684 |
| 190 | 1199 | 148 | 271 | 420 |
| 191 | 1200 | 1482 | 568 | 714 |
| 192 | 1201 | 1484 | 203 | 340 |
| 193 | 1202 | 1486 | 2185 | 1190 |
| 194 | 1203 | 1492 | 438 | 2912 |
| 195 | 1204 | 1493 | 82 | 225 |
| 196 | 1205 | 1501 | 210 | 347 |
| 197 | 1206 | 1508 | 1364 | 1101 |
| 198 | 1207 | 1509 | 56 | 613 |
| 199 | 1208 | 1512 | 828 | 965 |
| 200 | 1209 | 1515 | 3216 | 3812 |

# TABLE 3

| SEQ ID NO: OF NUCLEOTIDE | SEQ ID NO: OF AMINO ACID | SEQ ID NO: IN USSN 09/491,404 | START NUCLEOTIDE OF CODING REGION | STOP NUCLEOTIDE OF CODING REGION |
|---|---|---|---|---|
| 201 | 1210 | 1516 | 614 | 790 |
| 202 | 1211 | 1522 | 1709 | 1029 |
| 203 | 1212 | 1524 | 614 | 799 |
| 204 | 1213 | 1526 | 3917 | 4081 |
| 205 | 1214 | 1529 | 221 | 2146 |
| 206 | 1215 | 1530 | 644 | 390 |
| 207 | 1216 | 1532 | 16 | 1224 |
| 208 | 1217 | 1535 | 885 | 1031 |
| 209 | 1218 | 1536 | 245 | 1156 |
| 210 | 1219 | 1538 | 1617 | 4994 |
| 211 | 1220 | 154 | 97 | 234 |
| 212 | 1221 | 1540 | 4325 | 4158 |
| 213 | 1222 | 1541 | 2020 | 2778 |
| 214 | 1223 | 1544 | 595 | 3168 |
| 215 | 1224 | 1545 | 328 | 534 |
| 216 | 1225 | 1548 | 47 | 211 |
| 217 | 1226 | 1550 | 49 | 201 |
| 218 | 1227 | 1552 | 418 | 558 |
| 219 | 1228 | 1555 | 509 | 330 |
| 220 | 1229 | 1557 | 699 | 854 |
| 221 | 1230 | 1561 | 847 | 1932 |
| 222 | 1231 | 1563 | 775 | 933 |
| 223 | 1232 | 1565 | 286 | 453 |
| 224 | 1233 | 1567 | 807 | 974 |
| 225 | 1234 | 1568 | 1227 | 1601 |
| 226 | 1235 | 1569 | 113 | 328 |
| 227 | 1236 | 157 | 145 | 2 |
| 228 | 1237 | 1570 | 222 | 845 |
| 229 | 1238 | 1572 | 167 | 685 |
| 230 | 1239 | 1574 | 97 | 1167 |
| 231 | 1240 | 1575 | 581 | 2701 |
| 232 | 1241 | 1577 | 1246 | 953 |
| 233 | 1242 | 1578 | 1440 | 175 |
| 234 | 1243 | 1579 | 4738 | 4601 |
| 235 | 1244 | 1580 | 1431 | 1568 |
| 236 | 1245 | 1581 | 2491 | 3222 |
| 237 | 1246 | 1584 | 463 | 2157 |
| 238 | 1247 | 1585 | 156 | 2366 |
| 239 | 1248 | 1586 | 167 | 691 |
| 240 | 1249 | 1587 | 102 | 305 |
| 241 | 1250 | 1589 | 1157 | 1783 |
| 242 | 1251 | 159 | 812 | 639 |
| 243 | 1252 | 1592 | 270 | 521 |
| 244 | 1253 | 1593 | 92 | 310 |
| 245 | 1254 | 1594 | 814 | 188 |
| 246 | 1255 | 1595 | 101 | 2290 |
| 247 | 1256 | 1597 | 119 | 910 |
| 248 | 1257 | 1598 | 178 | 1398 |
| 249 | 1258 | 1600 | 2937 | 2578 |
| 250 | 1259 | 1604 | 47 | 526 |

## TABLE 3

| SEQ ID NO: OF NUCLEOTIDE | SEQ ID NO: OF AMINO ACID | SEQ ID NO: IN USSN 09/491,404 | START NUCLEOTIDE OF CODING REGION | STOP NUCLEOTIDE OF CODING REGION |
|---|---|---|---|---|
| 251 | 1260 | 1606 | 2204 | 1872 |
| 252 | 1261 | 1608 | 235 | 603 |
| 253 | 1262 | 1609 | 156 | 2366 |
| 254 | 1263 | 1611 | 1992 | 2135 |
| 255 | 1264 | 1614 | 968 | 786 |
| 256 | 1265 | 1615 | 2578 | 2751 |
| 257 | 1266 | 1616 | 6256 | 5813 |
| 258 | 1267 | 1617 | 29 | 709 |
| 259 | 1268 | 1619 | 1123 | 4071 |
| 260 | 1269 | 1621 | 581 | 2704 |
| 261 | 1270 | 1626 | 43 | 321 |
| 262 | 1271 | 1629 | 3616 | 1673 |
| 263 | 1272 | 163 | 509 | 183 |
| 264 | 1273 | 1630 | 81 | 248 |
| 265 | 1274 | 1631 | 9 | 572 |
| 266 | 1275 | 1633 | 2565 | 2807 |
| 267 | 1276 | 1634 | 2373 | 2510 |
| 268 | 1277 | 1635 | 3216 | 4508 |
| 269 | 1278 | 1636 | 4239 | 4081 |
| 270 | 1279 | 1642 | 4238 | 4020 |
| 271 | 1280 | 1643 | 152 | 304 |
| 272 | 1281 | 1644 | 47 | 478 |
| 273 | 1282 | 1645 | 121 | 921 |
| 274 | 1283 | 1646 | 3815 | 3030 |
| 275 | 1284 | 1647 | 335 | 186 |
| 276 | 1285 | 1649 | 6 | 974 |
| 277 | 1286 | 1654 | 34 | 951 |
| 278 | 1287 | 1655 | 491 | 1387 |
| 279 | 1288 | 1656 | 78 | 560 |
| 280 | 1289 | 1657 | 1431 | 1568 |
| 281 | 1290 | 1658 | 2373 | 1015 |
| 282 | 1291 | 1670 | 236 | 3 |
| 283 | 1292 | 1673 | 95 | 1342 |
| 284 | 1293 | 1685 | 2124 | 1786 |
| 285 | 1294 | 1690 | 245 | 415 |
| 286 | 1295 | 1691 | 977 | 774 |
| 287 | 1296 | 1699 | 50 | 247 |
| 288 | 1297 | 17 | 282 | 112 |
| 289 | 1298 | 1710 | 943 | 239 |
| 290 | 1299 | 1711 | 127 | 318 |
| 291 | 1300 | 1718 | 99 | 338 |
| 292 | 1301 | 1719 | 122 | 382 |
| 293 | 1302 | 172 | 33 | 461 |
| 294 | 1303 | 1720 | 180 | 1 |
| 295 | 1304 | 1722 | 160 | 327 |
| 296 | 1305 | 1726 | 175 | 363 |
| 297 | 1306 | 1737 | 84 | 497 |
| 298 | 1307 | 1738 | 188 | 379 |
| 299 | 1308 | 174 | 138 | 332 |
| 300 | 1309 | 1743 | 560 | 784 |

# TABLE 3

| SEQ ID NO: OF NUCLEOTIDE | SEQ ID NO: OF AMINO ACID | SEQ ID NO: IN USSN 09/491,404 | START NUCLEOTIDE OF CODING REGION | STOP NUCLEOTIDE OF CODING REGION |
|---|---|---|---|---|
| 301 | 1310 | 1747 | 1824 | 1961 |
| 302 | 1311 | 1748 | 97 | 411 |
| 303 | 1312 | 1749 | 151 | 492 |
| 304 | 1313 | 177 | 59 | 322 |
| 305 | 1314 | 1776 | 68 | 262 |
| 306 | 1315 | 1779 | 43 | 255 |
| 307 | 1316 | 178 | 58 | 399 |
| 308 | 1317 | 1781 | 1179 | 907 |
| 309 | 1318 | 1786 | 579 | 385 |
| 310 | 1319 | 1789 | 56 | 193 |
| 311 | 1320 | 180 | 218 | 78 |
| 312 | 1321 | 1800 | 230 | 394 |
| 313 | 1322 | 1801 | 1778 | 876 |
| 314 | 1323 | 181 | 174 | 428 |
| 315 | 1324 | 1829 | 179 | 42 |
| 316 | 1325 | 1846 | 525 | 785 |
| 317 | 1326 | 1848 | 5632 | 5838 |
| 318 | 1327 | 185 | 92 | 400 |
| 319 | 1328 | 1850 | 178 | 333 |
| 320 | 1329 | 186 | 699 | 1310 |
| 321 | 1330 | 1860 | 8 | 604 |
| 322 | 1331 | 1868 | 376 | 618 |
| 323 | 1332 | 187 | 148 | 366 |
| 324 | 1333 | 1870 | 233 | 388 |
| 325 | 1334 | 1872 | 12 | 206 |
| 326 | 1335 | 188 | 181 | 516 |
| 327 | 1336 | 1884 | 549 | 863 |
| 328 | 1337 | 1886 | 128 | 298 |
| 329 | 1338 | 189 | 28 | 204 |
| 330 | 1339 | 1891 | 11246 | 11097 |
| 331 | 1340 | 1895 | 175 | 417 |
| 332 | 1341 | 1897 | 221 | 400 |
| 333 | 1342 | 1899 | 744 | 890 |
| 334 | 1343 | 191 | 77 | 286 |
| 335 | 1344 | 1914 | 403 | 699 |
| 336 | 1345 | 192 | 8 | 343 |
| 337 | 1346 | 1947 | 656 | 1735 |
| 338 | 1347 | 1948 | 32 | 283 |
| 339 | 1348 | 195 | 129 | 323 |
| 340 | 1349 | 196 | 122 | 295 |
| 341 | 1350 | 1962 | 554 | 733 |
| 342 | 1351 | 197 | 110 | 277 |
| 343 | 1352 | 1976 | 348 | 2450 |
| 344 | 1353 | 198 | 93 | 239 |
| 345 | 1354 | 1980 | 137 | 310 |
| 346 | 1355 | 2 | 916 | 13698 |
| 347 | 1356 | 20 | 112 | 303 |
| 348 | 1357 | 2005 | 88 | 420 |
| 349 | 1358 | 2007 | 525 | 385 |
| 350 | 1359 | 2008 | 266 | 484 |

# TABLE 3

| SEQ ID NO: OF NUCLEOTIDE | SEQ ID NO: OF AMINO ACID | SEQ ID NO: IN USSN 09/491,404 | START NUCLEOTIDE OF CODING REGION | STOP NUCLEOTIDE OF CODING REGION |
|---|---|---|---|---|
| 351 | 1360 | 2013 | 64 | 234 |
| 352 | 1361 | 2016 | 99 | 329 |
| 353 | 1362 | 2018 | 84 | 401 |
| 354 | 1363 | 202 | 300 | 130 |
| 355 | 1364 | 2022 | 1240 | 1016 |
| 356 | 1365 | 2029 | 191 | 364 |
| 357 | 1366 | 2037 | 231 | 404 |
| 358 | 1367 | 2043 | 3206 | 3349 |
| 359 | 1368 | 2047 | 169 | 456 |
| 360 | 1369 | 2048 | 295 | 522 |
| 361 | 1370 | 2049 | 533 | 769 |
| 362 | 1371 | 205 | 4 | 684 |
| 363 | 1372 | 2051 | 403 | 699 |
| 364 | 1373 | 2055 | 173 | 379 |
| 365 | 1374 | 2056 | 270 | 1157 |
| 366 | 1375 | 2061 | 949 | 725 |
| 367 | 1376 | 2064 | 127 | 309 |
| 368 | 1377 | 2065 | 248 | 577 |
| 369 | 1378 | 2070 | 204 | 344 |
| 370 | 1379 | 2071 | 374 | 793 |
| 371 | 1380 | 2074 | 945 | 796 |
| 372 | 1381 | 2076 | 300 | 67 |
| 373 | 1382 | 2078 | 416 | 586 |
| 374 | 1383 | 2081 | 316 | 507 |
| 375 | 1384 | 2082 | 20 | 220 |
| 376 | 1385 | 209 | 19 | 168 |
| 377 | 1386 | 210 | 27 | 395 |
| 378 | 1387 | 2102 | 258 | 452 |
| 379 | 1388 | 2104 | 1706 | 1539 |
| 380 | 1389 | 211 | 84 | 311 |
| 381 | 1390 | 212 | 677 | 231 |
| 382 | 1391 | 2120 | 40 | 414 |
| 383 | 1392 | 214 | 101 | 268 |
| 384 | 1393 | 2140 | 213 | 377 |
| 385 | 1394 | 2161 | 216 | 368 |
| 386 | 1395 | 2162 | 106 | 420 |
| 387 | 1396 | 2164 | 104 | 250 |
| 388 | 1397 | 217 | 333 | 22 |
| 389 | 1398 | 218 | 80 | 325 |
| 390 | 1399 | 219 | 709 | 506 |
| 391 | 1400 | 2196 | 158 | 319 |
| 392 | 1401 | 2198 | 469 | 1164 |
| 393 | 1402 | 22 | 843 | 700 |
| 394 | 1403 | 2214 | 980 | 822 |
| 395 | 1404 | 2215 | 49 | 318 |
| 396 | 1405 | 2225 | 544 | 1974 |
| 397 | 1406 | 223 | 185 | 21 |
| 398 | 1407 | 2233 | 116 | 313 |
| 399 | 1408 | 224 | 189 | 16 |
| 400 | 1409 | 2240 | 2740 | 2525 |

# TABLE 3

| SEQ ID NO: OF NUCLEOTIDE | SEQ ID NO: OF AMINO ACID | SEQ ID NO: IN USSN 09/491,404 | START NUCLEOTIDE OF CODING REGION | STOP NUCLEOTIDE OF CODING REGION |
|---|---|---|---|---|
| 401 | 1410 | 2244 | 1489 | 1647 |
| 402 | 1411 | 2254 | 72 | 317 |
| 403 | 1412 | 226 | 335 | 120 |
| 404 | 1413 | 2260 | 562 | 738 |
| 405 | 1414 | 2268 | 300 | 67 |
| 406 | 1415 | 227 | 103 | 615 |
| 407 | 1416 | 2273 | 114 | 344 |
| 408 | 1417 | 2275 | 239 | 985 |
| 409 | 1418 | 2276 | 1358 | 1164 |
| 410 | 1419 | 2288 | 56 | 1459 |
| 411 | 1420 | 2291 | 83 | 532 |
| 412 | 1421 | 2296 | 264 | 530 |
| 413 | 1422 | 2298 | 533 | 781 |
| 414 | 1423 | 2300 | 1684 | 1845 |
| 415 | 1424 | 2305 | 8 | 226 |
| 416 | 1425 | 231 | 86 | 820 |
| 417 | 1426 | 232 | 361 | 1920 |
| 418 | 1427 | 233 | 150 | 467 |
| 419 | 1428 | 2331 | 334 | 2856 |
| 420 | 1429 | 2334 | 168 | 953 |
| 421 | 1430 | 2341 | 198 | 395 |
| 422 | 1431 | 2344 | 122 | 1432 |
| 423 | 1432 | 2346 | 1345 | 1187 |
| 424 | 1433 | 2348 | 502 | 729 |
| 425 | 1434 | 235 | 338 | 844 |
| 426 | 1435 | 2351 | 228 | 713 |
| 427 | 1436 | 236 | 232 | 2 |
| 428 | 1437 | 2360 | 1611 | 1357 |
| 429 | 1438 | 2362 | 36 | 263 |
| 430 | 1439 | 2364 | 294 | 1568 |
| 431 | 1440 | 2365 | 103 | 312 |
| 432 | 1441 | 2378 | 209 | 5281 |
| 433 | 1442 | 238 | 53 | 511 |
| 434 | 1443 | 2380 | 207 | 380 |
| 435 | 1444 | 239 | 457 | 663 |
| 436 | 1445 | 2392 | 176 | 2653 |
| 437 | 1446 | 2399 | 940 | 2040 |
| 438 | 1447 | 2405 | 144 | 380 |
| 439 | 1448 | 2407 | 1875 | 2702 |
| 440 | 1449 | 2415 | 1927 | 137 |
| 441 | 1450 | 242 | 1813 | 986 |
| 442 | 1451 | 2421 | 43 | 405 |
| 443 | 1452 | 2423 | 1556 | 1413 |
| 444 | 1453 | 2424 | 673 | 1041 |
| 445 | 1454 | 2432 | 295 | 1275 |
| 446 | 1455 | 2438 | 607 | 437 |
| 447 | 1456 | 2444 | 294 | 437 |
| 448 | 1457 | 2447 | 212 | 1588 |
| 449 | 1458 | 2448 | 52 | 1440 |
| 450 | 1459 | 2449 | 637 | 1197 |

# TABLE 3

| SEQ ID NO: OF NUCLEOTIDE | SEQ ID NO: OF AMINO ACID | SEQ ID NO: IN USSN 09/491,404 | START NUCLEOTIDE OF CODING REGION | STOP NUCLEOTIDE OF CODING REGION |
|---|---|---|---|---|
| 451 | 1460 | 245 | 208 | 876 |
| 452 | 1461 | 2450 | 3740 | 4369 |
| 453 | 1462 | 2453 | 222 | 389 |
| 454 | 1463 | 246 | 566 | 763 |
| 455 | 1464 | 2466 | 179 | 778 |
| 456 | 1465 | 2471 | 532 | 669 |
| 457 | 1466 | 2473 | 817 | 650 |
| 458 | 1467 | 2474 | 236 | 1333 |
| 459 | 1468 | 2476 | 173 | 3 |
| 460 | 1469 | 248 | 331 | 2 |
| 461 | 1470 | 2486 | 709 | 885 |
| 462 | 1471 | 249 | 88 | 456 |
| 463 | 1472 | 2496 | 107 | 1054 |
| 464 | 1473 | 2498 | 413 | 607 |
| 465 | 1474 | 2501 | 103 | 267 |
| 466 | 1475 | 2503 | 334 | 717 |
| 467 | 1476 | 2506 | 3740 | 4369 |
| 468 | 1477 | 2509 | 188 | 18 |
| 469 | 1478 | 2512 | 78 | 368 |
| 470 | 1479 | 2514 | 16 | 354 |
| 471 | 1480 | 2523 | 53 | 325 |
| 472 | 1481 | 2526 | 223 | 384 |
| 473 | 1482 | 2532 | 596 | 763 |
| 474 | 1483 | 2533 | 62 | 667 |
| 475 | 1484 | 2535 | 89 | 1519 |
| 476 | 1485 | 2537 | 175 | 375 |
| 477 | 1486 | 254 | 299 | 21 |
| 478 | 1487 | 2540 | 553 | 816 |
| 479 | 1488 | 2546 | 1905 | 1102 |
| 480 | 1489 | 2555 | 2046 | 4541 |
| 481 | 1490 | 2559 | 569 | 733 |
| 482 | 1491 | 256 | 9 | 410 |
| 483 | 1492 | 2560 | 288 | 76 |
| 484 | 1493 | 2565 | 3269 | 3502 |
| 485 | 1494 | 2569 | 116 | 478 |
| 486 | 1495 | 257 | 203 | 475 |
| 487 | 1496 | 2571 | 2763 | 2548 |
| 488 | 1497 | 2572 | 65 | 652 |
| 489 | 1498 | 2575 | 70 | 294 |
| 490 | 1499 | 2576 | 1195 | 1010 |
| 491 | 1500 | 258 | 434 | 21 |
| 492 | 1501 | 2580 | 155 | 400 |
| 493 | 1502 | 2591 | 53 | 214 |
| 494 | 1503 | 2592 | 163 | 348 |
| 495 | 1504 | 26 | 261 | 398 |
| 496 | 1505 | 2605 | 277 | 420 |
| 497 | 1506 | 261 | 29 | 598 |
| 498 | 1507 | 2614 | 1331 | 1510 |
| 499 | 1508 | 2617 | 235 | 378 |
| 500 | 1509 | 262 | 204 | 458 |

# TABLE 3

| SEQ ID NO: OF NUCLEOTIDE | SEQ ID NO: OF AMINO ACID | SEQ ID NO: IN USSN 09/491,404 | START NUCLEOTIDE OF CODING REGION | STOP NUCLEOTIDE OF CODING REGION |
|---|---|---|---|---|
| 501 | 1510 | 2624 | 254 | 418 |
| 502 | 1511 | 263 | 247 | 570 |
| 503 | 1512 | 264 | 184 | 540 |
| 504 | 1513 | 2643 | 1108 | 4026 |
| 505 | 1514 | 2644 | 305 | 535 |
| 506 | 1515 | 2645 | 1952 | 1509 |
| 507 | 1516 | 2647 | 1225 | 404 |
| 508 | 1517 | 2648 | 41 | 778 |
| 509 | 1518 | 265 | 53 | 418 |
| 510 | 1519 | 2650 | 190 | 936 |
| 511 | 1520 | 2658 | 1576 | 2451 |
| 512 | 1521 | 2659 | 44 | 430 |
| 513 | 1522 | 266 | 350 | 153 |
| 514 | 1523 | 2663 | 785 | 1177 |
| 515 | 1524 | 2665 | 395 | 550 |
| 516 | 1525 | 2666 | 41 | 778 |
| 517 | 1526 | 2667 | 244 | 384 |
| 518 | 1527 | 2668 | 174 | 527 |
| 519 | 1528 | 2669 | 27 | 302 |
| 520 | 1529 | 2678 | 1172 | 960 |
| 521 | 1530 | 2684 | 178 | 432 |
| 522 | 1531 | 269 | 341 | 520 |
| 523 | 1532 | 2699 | 1241 | 1083 |
| 524 | 1533 | 2701 | 402 | 2624 |
| 525 | 1534 | 2702 | 28 | 177 |
| 526 | 1535 | 2706 | 1108 | 4026 |
| 527 | 1536 | 2707 | 1240 | 1016 |
| 528 | 1537 | 271 | 59 | 346 |
| 529 | 1538 | 2714 | 34 | 987 |
| 530 | 1539 | 2715 | 1117 | 647 |
| 531 | 1540 | 2717 | 25 | 429 |
| 532 | 1541 | 2718 | 1670 | 1885 |
| 533 | 1542 | 2719 | 31 | 1137 |
| 534 | 1543 | 272 | 6 | 152 |
| 535 | 1544 | 2726 | 230 | 592 |
| 536 | 1545 | 2728 | 578 | 369 |
| 537 | 1546 | 2731 | 193 | 366 |
| 538 | 1547 | 2735 | 495 | 301 |
| 539 | 1548 | 274 | 352 | 119 |
| 540 | 1549 | 2741 | 94 | 255 |
| 541 | 1550 | 2798 | 1031 | 1240 |
| 542 | 1551 | 28 | 54 | 725 |
| 543 | 1552 | 2803 | 204 | 374 |
| 544 | 1553 | 2809 | 216 | 938 |
| 545 | 1554 | 2822 | 280 | 447 |
| 546 | 1555 | 2823 | 197 | 388 |
| 547 | 1556 | 2824 | 224 | 12 |
| 548 | 1557 | 2826 | 79 | 456 |
| 549 | 1558 | 2828 | 24 | 428 |
| 550 | 1559 | 2838 | 90 | 698 |

# TABLE 3

| SEQ ID NO: OF NUCLEOTIDE | SEQ ID NO: OF AMINO ACID | SEQ ID NO: IN USSN 09/491,404 | START NUCLEOTIDE OF CODING REGION | STOP NUCLEOTIDE OF CODING REGION |
|---|---|---|---|---|
| 551 | 1560 | 284 | 21 | 197 |
| 552 | 1561 | 2847 | 113 | 262 |
| 553 | 1562 | 285 | 146 | 292 |
| 554 | 1563 | 2852 | 233 | 439 |
| 555 | 1564 | 2854 | 830 | 988 |
| 556 | 1565 | 2855 | 336 | 1043 |
| 557 | 1566 | 2856 | 384 | 614 |
| 558 | 1567 | 2857 | 437 | 748 |
| 559 | 1568 | 2859 | 1295 | 1158 |
| 560 | 1569 | 286 | 30 | 179 |
| 561 | 1570 | 2860 | 2618 | 2469 |
| 562 | 1571 | 2864 | 1325 | 1176 |
| 563 | 1572 | 2867 | 1034 | 795 |
| 564 | 1573 | 288 | 190 | 345 |
| 565 | 1574 | 2884 | 856 | 257 |
| 566 | 1575 | 2886 | 15 | 167 |
| 567 | 1576 | 2891 | 34 | 405 |
| 568 | 1577 | 2900 | 104 | 2683 |
| 569 | 1578 | 2901 | 193 | 366 |
| 570 | 1579 | 2902 | 91 | 1806 |
| 571 | 1580 | 2907 | 268 | 498 |
| 572 | 1581 | 2908 | 83 | 1564 |
| 573 | 1582 | 2910 | 2131 | 3117 |
| 574 | 1583 | 2915 | 715 | 861 |
| 575 | 1584 | 2916 | 52 | 2064 |
| 576 | 1585 | 2919 | 62 | 1015 |
| 577 | 1586 | 292 | 615 | 854 |
| 578 | 1587 | 2923 | 332 | 1279 |
| 579 | 1588 | 2924 | 264 | 422 |
| 580 | 1589 | 2925 | 122 | 1432 |
| 581 | 1590 | 2930 | 195 | 341 |
| 582 | 1591 | 2931 | 221 | 3 |
| 583 | 1592 | 2934 | 1642 | 1827 |
| 584 | 1593 | 2937 | 38 | 421 |
| 585 | 1594 | 2940 | 520 | 383 |
| 586 | 1595 | 2944 | 325 | 68 |
| 587 | 1596 | 295 | 49 | 255 |
| 588 | 1597 | 2950 | 226 | 59 |
| 589 | 1598 | 2951 | 110 | 400 |
| 590 | 1599 | 2955 | 303 | 641 |
| 591 | 1600 | 2957 | 365 | 673 |
| 592 | 1601 | 2964 | 96 | 347 |
| 593 | 1602 | 2967 | 738 | 466 |
| 594 | 1603 | 2968 | 222 | 428 |
| 595 | 1604 | 2969 | 365 | 117 |
| 596 | 1605 | 2970 | 314 | 643 |
| 597 | 1606 | 2973 | 961 | 1176 |
| 598 | 1607 | 2975 | 975 | 799 |
| 599 | 1608 | 2979 | 89 | 442 |
| 600 | 1609 | 298 | 152 | 3 |

# TABLE 3

| SEQ ID NO: OF NUCLEOTIDE | SEQ ID NO: OF AMINO ACID | SEQ ID NO: IN USSN 09/491,404 | START NUCLEOTIDE OF CODING REGION | STOP NUCLEOTIDE OF CODING REGION |
|---|---|---|---|---|
| 601 | 1610 | 2991 | 112 | 261 |
| 602 | 1611 | 2995 | 201 | 368 |
| 603 | 1612 | 3 | 13559 | 13335 |
| 604 | 1613 | 30 | 176 | 751 |
| 605 | 1614 | 3002 | 1807 | 2265 |
| 606 | 1615 | 3005 | 339 | 743 |
| 607 | 1616 | 3023 | 64 | 243 |
| 608 | 1617 | 3039 | 71 | 217 |
| 609 | 1618 | 304 | 50 | 334 |
| 610 | 1619 | 305 | 226 | 387 |
| 611 | 1620 | 3051 | 56 | 268 |
| 612 | 1621 | 307 | 9. | 278 |
| 613 | 1622 | 308 | 116 | 274 |
| 614 | 1623 | 3085 | 97 | 3030 |
| 615 | 1624 | 3088 | 801 | 634 |
| 616 | 1625 | 3089 | 18 | 455 |
| 617 | 1626 | 3094 | 92 | 1246 |
| 618 | 1627 | 3098 | 40 | 342 |
| 619 | 1628 | 310 | 142 | 354 |
| 620 | 1629 | 3101 | 48 | 383 |
| 621 | 1630 | 3105 | 188 | 328 |
| 622 | 1631 | 3107 | 177 | 413 |
| 623 | 1632 | 3109 | 184 | 327 |
| 624 | 1633 | 3114 | 70 | 243 |
| 625 | 1634 | 3115 | 295 | 459 |
| 626 | 1635 | 3116 | 115 | 348 |
| 627 | 1636 | 3119 | 70 | 222 |
| 628 | 1637 | 3120 | 163 | 531 |
| 629 | 1638 | 3122 | 60 | 266 |
| 630 | 1639 | 3129 | 226 | 501 |
| 631 | 1640 | 3146 | 190 | 363 |
| 632 | 1641 | 3151 | 212 | 1588 |
| 633 | 1642 | 3153 | 86 | 517 |
| 634 | 1643 | 3165 | 244 | 453 |
| 635 | 1644 | 317 | 97 | 342 |
| 636 | 1645 | 3179 | 106 | 873 |
| 637 | 1646 | 3181 | 108 | 896 |
| 638 | 1647 | 3182 | 554 | 775 |
| 639 | 1648 | 3192 | 268 | 441 |
| 640 | 1649 | 3194 | 923 | 1192 |
| 641 | 1650 | 3195 | 38 | 376 |
| 642 | 1651 | 32 | 185 | 334 |
| 643 | 1652 | 3200 | 199 | 561 |
| 644 | 1653 | 3201 | 516 | 848 |
| 645 | 1654 | 3202 | 232 | 681 |
| 646 | 1655 | 3208 | 836 | 633 |
| 647 | 1656 | 3210 | 202 | 384 |
| 648 | 1657 | 3214 | 349 | 588 |
| 649 | 1658 | 3215 | 859 | 380 |
| 650 | 1659 | 3216 | 51 | 320 |

# TABLE 3

| SEQ ID NO: OF NUCLEOTIDE | SEQ ID NO: OF AMINO ACID | SEQ ID NO: IN USSN 09/491,404 | START NUCLEOTIDE OF CODING REGION | STOP NUCLEOTIDE OF CODING REGION |
|---|---|---|---|---|
| 651 | 1660 | 3220 | 116 | 283 |
| 652 | 1661 | 3222 | 324 | 545 |
| 653 | 1662 | 3227 | 385 | 1197 |
| 654 | 1663 | 323 | 65 | 223 |
| 655 | 1664 | 3240 | 385 | 1197 |
| 656 | 1665 | 3243 | 65 | 916 |
| 657 | 1666 | 3250 | 263 | 463 |
| 658 | 1667 | 3252 | 244 | 480 |
| 659 | 1668 | 3253 | 136 | 297 |
| 660 | 1669 | 3254 | 83 | 439 |
| 661 | 1670 | 3255 | 573 | 920 |
| 662 | 1671 | 3257 | 548 | 757 |
| 663 | 1672 | 3259 | 34 | 822 |
| 664 | 1673 | 326 | 58 | 525 |
| 665 | 1674 | 3263 | 102 | 350 |
| 666 | 1675 | 3270 | 313 | 152 |
| 667 | 1676 | 3271 | 117 | 473 |
| 668 | 1677 | 3272 | 44 | 190 |
| 669 | 1678 | 3273 | 106 | 486 |
| 670 | 1679 | 3274 | 246 | 392 |
| 671 | 1680 | 3278 | 174 | 1 |
| 672 | 1681 | 3281 | 988 | 1134 |
| 673 | 1682 | 3282 | 101 | 334 |
| 674 | 1683 | 3291 | 129 | 284 |
| 675 | 1684 | 3294 | 101 | 595 |
| 676 | 1685 | 3296 | 107 | 565 |
| 677 | 1686 | 3298 | 130 | 552 |
| 678 | 1687 | 3299 | 333 | 515 |
| 679 | 1688 | 3300 | 324 | 121 |
| 680 | 1689 | 3303 | 378 | 157 |
| 681 | 1690 | 3306 | 296 | 637 |
| 682 | 1691 | 3307 | 1454 | 1660 |
| 683 | 1692 | 3309 | 163 | 471 |
| 684 | 1693 | 3311 | 335 | 478 |
| 685 | 1694 | 3312 | 5 | 280 |
| 686 | 1695 | 3313 | 298 | 546 |
| 687 | 1696 | 3314 | 50 | 526 |
| 688 | 1697 | 3315 | 99 | 413 |
| 689 | 1698 | 3322 | 101 | 685 |
| 690 | 1699 | 3323 | 66 | 356 |
| 691 | 1700 | 3324 | 76 | 462 |
| 692 | 1701 | 3328 | 248 | 904 |
| 693 | 1702 | 3335 | 136 | 393 |
| 694 | 1703 | 3336 | 47 | 733 |
| 695 | 1704 | 3338 | 181 | 786 |
| 696 | 1705 | 3339 | 58 | 231 |
| 697 | 1706 | 3342 | 226 | 390 |
| 698 | 1707 | 3349 | 72 | 488 |
| 699 | 1708 | 3356 | 208 | 384 |
| 700 | 1709 | 3358 | 194 | 436 |

## TABLE 3

| SEQ ID NO: OF NUCLEOTIDE | SEQ ID NO: OF AMINO ACID | SEQ ID NO: IN USSN 09/491,404 | START NUCLEOTIDE OF CODING REGION | STOP NUCLEOTIDE OF CODING REGION |
|---|---|---|---|---|
| 701 | 1710 | 3360 | 263 | 1459 |
| 702 | 1711 | 3366 | 55 | 816 |
| 703 | 1712 | 3367 | 364 | 735 |
| 704 | 1713 | 3370 | 237 | 878 |
| 705 | 1714 | 3371 | 188 | 721 |
| 706 | 1715 | 3372 | 14 | 241 |
| 707 | 1716 | 3373 | 42 | 290 |
| 708 | 1717 | 3387 | 32 | 202 |
| 709 | 1718 | 3389 | 29 | 256 |
| 710 | 1719 | 3390 | 181 | 393 |
| 711 | 1720 | 3396 | 520 | 822 |
| 712 | 1721 | 3410 | 10 | 153 |
| 713 | 1722 | 3412 | 82 | 291 |
| 714 | 1723 | 3414 | 453 | 292 |
| 715 | 1724 | 3421 | 158 | 337 |
| 716 | 1725 | 3427 | 430 | 618 |
| 717 | 1726 | 3430 | 210 | 380 |
| 718 | 1727 | 3431 | 295 | 432 |
| 719 | 1728 | 3440 | 419 | 556 |
| 720 | 1729 | 3444 | 402 | 256 |
| 721 | 1730 | 3445 | 281 | 430 |
| 722 | 1731 | 346 | 42 | 722 |
| 723 | 1732 | 347 | 384 | 689 |
| 724 | 1733 | 3470 | 114 | 530 |
| 725 | 1734 | 3478 | 38 | 217 |
| 726 | 1735 | 3479 | 161 | 379 |
| 727 | 1736 | 348 | 37 | 231 |
| 728 | 1737 | 3482 | 156 | 296 |
| 729 | 1738 | 35 | 255 | 575 |
| 730 | 1739 | 3503 | 185 | 454 |
| 731 | 1740 | 3505 | 252 | 422 |
| 732 | 1741 | 3529 | 37 | 183 |
| 733 | 1742 | 353 | 262 | 522 |
| 734 | 1743 | 3537 | 127 | 273 |
| 735 | 1744 | 3539 | 98 | 268 |
| 736 | 1745 | 3542 | 25 | 312 |
| 737 | 1746 | 3543 | 70 | 228 |
| 738 | 1747 | 3544 | 31 | 177 |
| 739 | 1748 | 3548 | 972 | 385 |
| 740 | 1749 | 3553 | 27 | 164 |
| 741 | 1750 | 3560 | 113 | 358 |
| 742 | 1751 | 3563 | 483 | 764 |
| 743 | 1752 | 3564 | 6 | 434 |
| 744 | 1753 | 3566 | 316 | 507 |
| 745 | 1754 | 3570 | 6 | 377 |
| 746 | 1755 | 3574 | 108 | 440 |
| 747 | 1756 | 3576 | 569 | 348 |
| 748 | 1757 | 3579 | 293 | 442 |
| 749 | 1758 | 3582 | 20 | 388 |
| 750 | 1759 | 3583 | 172 | 396 |

# TABLE 3

| SEQ ID NO: OF NUCLEOTIDE | SEQ ID NO: OF AMINO ACID | SEQ ID NO: IN USSN 09/491,404 | START NUCLEOTIDE OF CODING REGION | STOP NUCLEOTIDE OF CODING REGION |
|---|---|---|---|---|
| 751 | 1760 | 3587 | 84 | 449 |
| 752 | 1761 | 3596 | 91 | 459 |
| 753 | 1762 | 3599 | 40 | 474 |
| 754 | 1763 | 3606 | 335 | 1105 |
| 755 | 1764 | 3609 | 169 | 666 |
| 756 | 1765 | 3617 | 141 | 410 |
| 757 | 1766 | 3620 | 218 | 388 |
| 758 | 1767 | 3630 | 189 | 1 |
| 759 | 1768 | 3642 | 122 | 643 |
| 760 | 1769 | 3644 | 431 | 664 |
| 761 | 1770 | 3647 | 274 | 720 |
| 762 | 1771 | 3651 | 245 | 472 |
| 763 | 1772 | 3652 | 259 | 642 |
| 764 | 1773 | 3653 | 153 | 1994 |
| 765 | 1774 | 3654 | 87 | 554 |
| 766 | 1775 | 3657 | 57 | 2744 |
| 767 | 1776 | 3658 | 387 | 920 |
| 768 | 1777 | 366 | 402 | 578 |
| 769 | 1778 | 3660 | 120 | 530 |
| 770 | 1779 | 3661 | 480 | 674 |
| 771 | 1780 | 3663 | 1096 | 938 |
| 772 | 1781 | 3669 | 689 | 1015 |
| 773 | 1782 | 3677 | 469 | 642 |
| 774 | 1783 | 3678 | 1194 | 889 |
| 775 | 1784 | 3685 | 406 | 1134 |
| 776 | 1785 | 3689 | 233 | 706 |
| 777 | 1786 | 3693 | 21 | 446 |
| 778 | 1787 | 3699 | 55 | 414 |
| 779 | 1788 | 370 | 59 | 262 |
| 780 | 1789 | 3707 | 38 | 436 |
| 781 | 1790 | 3711 | 229 | 474 |
| 782 | 1791 | 3713 | 314 | 463 |
| 783 | 1792 | 3717 | 178 | 675 |
| 784 | 1793 | 3720 | 258 | 695 |
| 785 | 1794 | 3721 | 96 | 548 |
| 786 | 1795 | 3722 | 32 | 562 |
| 787 | 1796 | 3724 | 220 | 513 |
| 788 | 1797 | 3726 | 180 | 467 |
| 789 | 1798 | 3729 | 251 | 523 |
| 790 | 1799 | 373 | 110 | 340 |
| 791 | 1800 | 3735 | 91 | 636 |
| 792 | 1801 | 3736 | 275 | 880 |
| 793 | 1802 | 3738 | 106 | 621 |
| 794 | 1803 | 3762 | 702 | 1175 |
| 795 | 1804 | 3768 | 293 | 598 |
| 796 | 1805 | 377 | 96 | 257 |
| 797 | 1806 | 3772 | 169 | 2 |
| 798 | 1807 | 3786 | 108 | 248 |
| 799 | 1808 | 3787 | 282 | 638 |
| 800 | 1809 | 3789 | 139 | 411 |

# TABLE 3

| SEQ ID NO: OF NUCLEOTIDE | SEQ ID NO: OF AMINO ACID | SEQ ID NO: IN USSN 09/491,404 | START NUCLEOTIDE OF CODING REGION | STOP NUCLEOTIDE OF CODING REGION |
|---|---|---|---|---|
| 801 | 1810 | 379 | 248 | 421 |
| 802 | 1811 | 38 | 146 | 3 |
| 803 | 1812 | 382 | 24 | 275 |
| 804 | 1813 | 385 | 138 | 1 |
| 805 | 1814 | 388 | 268 | 74 |
| 806 | 1815 | 39 | 302 | 3 |
| 807 | 1816 | 391 | 24 | 368 |
| 808 | 1817 | 395 | 51 | 482 |
| 809 | 1818 | 397 | 422 | 766 |
| 810 | 1819 | 399 | 102 | 311 |
| 811 | 1820 | 4 | 11219 | 13123 |
| 812 | 1821 | 405 | 253 | 2 |
| 813 | 1822 | 406 | 342 | 665 |
| 814 | 1823 | 411 | 321 | 542 |
| 815 | 1824 | 416 | 736 | 909 |
| 816 | 1825 | 422 | 1541 | 867 |
| 817 | 1826 | 43 | 330 | 686 |
| 818 | 1827 | 434 | 207 | 34 |
| 819 | 1828 | 435 | 140 | 445 |
| 820 | 1829 | 437 | 160 | 423 |
| 821 | 1830 | 439 | 347 | 706 |
| 822 | 1831 | 44 | 91 | 282 |
| 823 | 1832 | 450 | 136 | 402 |
| 824 | 1833 | 458 | 169 | 348 |
| 825 | 1834 | 459 | 99 | 284 |
| 826 | 1835 | 462 | 70 | 282 |
| 827 | 1836 | 465 | 462 | 791 |
| 828 | 1837 | 467 | 76 | 348 |
| 829 | 1838 | 470 | 35 | 637 |
| 830 | 1839 | 475 | 37 | 426 |
| 831 | 1840 | 477 | 242 | 382 |
| 832 | 1841 | 478 | 66 | 311 |
| 833 | 1842 | 485 | 196 | 426 |
| 834 | 1843 | 488 | 117 | 443 |
| 835 | 1844 | 490 | 231 | 485 |
| 836 | 1845 | 493 | 281 | 610 |
| 837 | 1846 | 496 | 90 | 371 |
| 838 | 1847 | 5 | 34 | 3933 |
| 839 | 1848 | 501 | 60 | 368 |
| 840 | 1849 | 502 | 707 | 856 |
| 841 | 1850 | 504 | 208 | 459 |
| 842 | 1851 | 505 | 165 | 317 |
| 843 | 1852 | 509 | 62 | 223 |
| 844 | 1853 | 511 | 46 | 432 |
| 845 | 1854 | 515 | 13 | 582 |
| 846 | 1855 | 516 | 92 | 325 |
| 847 | 1856 | 518 | 83 | 283 |
| 848 | 1857 | 519 | 365 | 685 |
| 849 | 1858 | 521 | 12 | 413 |
| 850 | 1859 | 525 | 6 | 251 |

# TABLE 3

| SEQ ID NO: OF NUCLEOTIDE | SEQ ID NO: OF AMINO ACID | SEQ ID NO: IN USSN 09/491,404 | START NUCLEOTIDE OF CODING REGION | STOP NUCLEOTIDE OF CODING REGION |
|---|---|---|---|---|
| 851 | 1860 | 526 | 862 | 725 |
| 852 | 1861 | 532 | 207 | 590 |
| 853 | 1862 | 536 | 226 | 53 |
| 854 | 1863 | 537 | 49 | 198 |
| 855 | 1864 | 540 | 270 | 1 |
| 856 | 1865 | 541 | 38 | 412 |
| 857 | 1866 | 546 | 388 | 2 |
| 858 | 1867 | 555 | 199 | 438 |
| 859 | 1868 | 556 | 144 | 482 |
| 860 | 1869 | 559 | 380 | 165 |
| 861 | 1870 | 563 | 27 | 617 |
| 862 | 1871 | 566 | 158 | 382 |
| 863 | 1872 | 568 | 69 | 320 |
| 864 | 1873 | 57 | 6 | 158 |
| 865 | 1874 | 571 | 8 | 1516 |
| 866 | 1875 | 572 | 32 | 505 |
| 867 | 1876 | 573 | 139 | 456 |
| 868 | 1877 | 574 | 49 | 771 |
| 869 | 1878 | 576 | 519 | 370 |
| 870 | 1879 | 578 | 168 | 1 |
| 871 | 1880 | 580 | 159 | 641 |
| 872 | 1881 | 581 | 108 | 497 |
| 873 | 1882 | 582 | 80 | 403 |
| 874 | 1883 | 587 | 172 | 435 |
| 875 | 1884 | 589 | 27 | 374 |
| 876 | 1885 | 590 | 84 | 428 |
| 877 | 1886 | 595 | 68 | 1138 |
| 878 | 1887 | 598 | 1023 | 766 |
| 879 | 1888 | 61 | 65 | 208 |
| 880 | 1889 | 612 | 310 | 546 |
| 881 | 1890 | 614 | 166 | 918 |
| 882 | 1891 | 617 | 252 | 602 |
| 883 | 1892 | 62 | 969 | 661 |
| 884 | 1893 | 620 | 188 | 418 |
| 885 | 1894 | 622 | 877 | 1014 |
| 886 | 1895 | 629 | 202 | 687 |
| 887 | 1896 | 63 | 98 | 277 |
| 888 | 1897 | 632 | 221 | 367 |
| 889 | 1898 | 64 | 536 | 381 |
| 890 | 1899 | 640 | 338 | 3 |
| 891 | 1900 | 641 | 12 | 395 |
| 892 | 1901 | 642 | 194 | 397 |
| 893 | 1902 | 644 | 15 | 395 |
| 894 | 1903 | 646 | 132 | 380 |
| 895 | 1904 | 647 | 3 | 389 |
| 896 | 1905 | 650 | 135 | 413 |
| 897 | 1906 | 651 | 231 | 428 |
| 898 | 1907 | 653 | 128 | 442 |
| 899 | 1908 | 654 | 214 | 77 |
| 900 | 1909 | 656 | 49 | 465 |

# TABLE 3

| SEQ ID NO: OF NUCLEOTIDE | SEQ ID NO: OF AMINO ACID | SEQ ID NO: IN USSN 09/491,404 | START NUCLEOTIDE OF CODING REGION | STOP NUCLEOTIDE OF CODING REGION |
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| 902 | 1911 | 66 | 267 | 614 |
| 903 | 1912 | 662 | 387 | 701 |
| 904 | 1913 | 666 | 76 | 498 |
| 905 | 1914 | 667 | 517 | 2184 |
| 906 | 1915 | 668 | 1423 | 788 |
| 907 | 1916 | 67 | 107 | 622 |
| 908 | 1917 | 678 | 172 | 387 |
| 909 | 1918 | 68 | 78 | 341 |
| 910 | 1919 | 680 | 832 | 671 |
| 911 | 1920 | 683 | 505 | 164 |
| 912 | 1921 | 687 | 105 | 521 |
| 913 | 1922 | 690 | 139 | 294 |
| 914 | 1923 | 691 | 244 | 456 |
| 915 | 1924 | 699 | 194 | 754 |
| 916 | 1925 | 701 | 371 | 520 |
| 917 | 1926 | 702 | 1888 | 2028 |
| 918 | 1927 | 704 | 1254 | 808 |
| 919 | 1928 | 705 | 126 | 1463 |
| 920 | 1929 | 706 | 31 | 390 |
| 921 | 1930 | 707 | 367 | 2 |
| 922 | 1931 | 709 | 1152 | 934 |
| 923 | 1932 | 715 | 744 | 541 |
| 924 | 1933 | 716 | 1360 | 1220 |
| 925 | 1934 | 722 | 173 | 430 |
| 926 | 1935 | 725 | 498 | 271 |
| 927 | 1936 | 727 | 18 | 164 |
| 928 | 1937 | 729 | 230 | 3 |
| 929 | 1938 | 73 | 262 | 834 |
| 930 | 1939 | 731 | 491 | 246 |
| 931 | 1940 | 740 | 20 | 322 |
| 932 | 1941 | 741 | 1430 | 1167 |
| 933 | 1942 | 747 | 660 | 523 |
| 934 | 1943 | 749 | 263 | 727 |
| 935 | 1944 | 750 | 209 | 391 |
| 936 | 1945 | 751 | 753 | 517 |
| 937 | 1946 | 755 | 172 | 387 |
| 938 | 1947 | 756 | 209 | 376 |
| 939 | 1948 | 76 | 656 | 513 |
| 940 | 1949 | 760 | 131 | 538 |
| 941 | 1950 | 763 | 893 | 1126 |
| 942 | 1951 | 766 | 1271 | 1537 |
| 943 | 1952 | 771 | 458 | 318 |
| 944 | 1953 | 775 | 391 | 558 |
| 945 | 1954 | 781 | 410 | 1684 |
| 946 | 1955 | 791 | 967 | 1284 |
| 947 | 1956 | 793 | 554 | 970 |
| 948 | 1957 | 795 | 8 | 268 |
| 949 | 1958 | 796 | 342 | 199 |
| 950 | 1959 | 798 | 211 | 405 |

# TABLE 3

| SEQ ID NO: OF NUCLEOTIDE | SEQ ID NO: OF AMINO ACID | SEQ ID NO: IN USSN 09/491,404 | START NUCLEOTIDE OF CODING REGION | STOP NUCLEOTIDE OF CODING REGION |
|---|---|---|---|---|
| 951 | 1960 | 799 | 625 | 392 |
| 952 | 1961 | 8 | 1523 | 1293 |
| 953 | 1962 | 801 | 484 | 678 |
| 954 | 1963 | 802 | 331 | 489 |
| 955 | 1964 | 808 | 210 | 905 |
| 956 | 1965 | 812 | 162 | 920 |
| 957 | 1966 | 819 | 723 | 2669 |
| 958 | 1967 | 820 | 964 | 725 |
| 959 | 1968 | 825 | 182 | 328 |
| 960 | 1969 | 829 | 1843 | 2292 |
| 961 | 1970 | 830 | 58 | 201 |
| 962 | 1971 | 832 | 150 | 341 |
| 963 | 1972 | 835 | 130 | 762 |
| 964 | 1973 | 836 | 449 | 291 |
| 965 | 1974 | 838 | 175 | 324 |
| 966 | 1975 | 84 | 175 | 435 |
| 967 | 1976 | 842 | 73 | 393 |
| 968 | 1977 | 844 | 423 | 824 |
| 969 | 1978 | 845 | 214 | 32 |
| 970 | 1979 | 846 | 120 | 317 |
| 971 | 1980 | 847 | 212 | 364 |
| 972 | 1981 | 85 | 190 | 426 |
| 973 | 1982 | 852 | 74 | 541 |
| 974 | 1983 | 855 | 1653 | 1465 |
| 975 | 1984 | 857 | 1964 | 2659 |
| 976 | 1985 | 858 | 598 | 1020 |
| 977 | 1986 | 861 | 58 | 933 |
| 978 | 1987 | 876 | 222 | 779 |
| 979 | 1988 | 878 | 2021 | 2161 |
| 980 | 1989 | 879 | 189 | 362 |
| 981 | 1990 | 88 | 39 | 278 |
| 982 | 1991 | 886 | 1165 | 1022 |
| 983 | 1992 | 891 | 158 | 310 |
| 984 | 1993 | 892 | 759 | 995 |
| 985 | 1994 | 895 | 224 | 379 |
| 986 | 1995 | 897 | 131 | 622 |
| 987 | 1996 | 9 | 1678 | 1448 |
| 988 | 1997 | 901 | 55 | 753 |
| 989 | 1998 | 906 | 450 | 623 |
| 990 | 1999 | 913 | 40 | 237 |
| 991 | 2000 | 918 | 17 | 334 |
| 992 | 2001 | 92 | 385 | 122 |
| 993 | 2002 | 926 | 772 | 518 |
| 994 | 2003 | 929 | 146 | 283 |
| 995 | 2004 | 932 | 23 | 175 |
| 996 | 2005 | 934 | 38 | 235 |
| 997 | 2006 | 935 | 286 | 423 |
| 998 | 2007 | 936 | 24 | 284 |
| 999 | 2008 | 939 | 450 | 623 |
| 1000 | 2009 | 94 | 139 | 2 |

## TABLE 3

| SEQ ID NO: OF NUCLEOTIDE | SEQ ID NO: OF AMINO ACID | SEQ ID NO: IN USSN 09/491,404 | START NUCLEOTIDE OF CODING REGION | STOP NUCLEOTIDE OF CODING REGION |
|---|---|---|---|---|
| 1001 | 2010 | 944 | 156 | 860 |
| 1002 | 2011 | 947 | 174 | 356 |
| 1003 | 2012 | 957 | 80 | 400 |
| 1004 | 2013 | 96 | 187 | 387 |
| 1005 | 2014 | 964 | 1352 | 1528 |
| 1006 | 2015 | 97 | 166 | 2 |
| 1007 | 2016 | 98 | 535 | 344 |
| 1008 | 2017 | 995 | 559 | 386 |
| 1009 | 2018 | 997 | 34 | 231 |

WHAT IS CLAIMED IS:

1. An isolated polynucleotide comprising a nucleotide sequence selected from the group consisting of SEQ ID NO: 1-1009, a mature protein coding portion of SEQ ID NO: 1-1009, an active domain of SEQ ID NO: 1-1009, and complementary sequences thereof.

2. An isolated polynucleotide encoding a polypeptide with biological activity, wherein said polynucleotide hybridizes to the polynucleotide of claim 1 under stringent hybridization conditions.

3. An isolated polynucleotide encoding a polypeptide with biological activity, wherein said polynucleotide has greater than about 90% sequence identity with the polynucleotide of claim 1.

4. The polynucleotide of claim 1 wherein said polynucleotide is DNA.

5. An isolated polynucleotide of claim 1 wherein said polynucleotide comprises the complementary sequences.

6. A vector comprising the polynucleotide of claim 1.

7. An expression vector comprising the polynucleotide of claim 1.

8. A host cell genetically engineered to comprise the polynucleotide of claim 1.

9. A host cell genetically engineered to comprise the polynucleotide of claim 1 operatively associated with a regulatory sequence that modulates expression of the polynucleotide in the host cell.

10. An isolated polypeptide, wherein the polypeptide is selected from the group consisting of:

(a) a polypeptide encoded by any one of the polynucleotides of claim 1; and

(b) a polypeptide encoded by a polynucleotide hybridizing under stringent conditions with any one of SEQ ID NO:1-1009.

11. A composition comprising the polypeptide of claim 10 and a carrier.

12. An antibody directed against the polypeptide of claim 10.

13. A method for detecting the polynucleotide of claim 1 in a sample, comprising:

a) contacting the sample with a compound that binds to and forms a complex with the polynucleotide of claim 1 for a period sufficient to form the complex; and

b) detecting the complex, so that if a complex is detected, the polynucleotide of claim 1 is detected.

14. A method for detecting the polynucleotide of claim 1 in a sample, comprising:

a) contacting the sample under stringent hybridization conditions with nucleic acid primers that anneal to the polynucleotide of claim 1 under such conditions;

b) amplifying a product comprising at least a portion of the polynucleotide of claim 1; and

c) detecting said product and thereby the polynucleotide of claim 1 in the sample.

15. The method of claim 14, wherein the polynucleotide is an RNA molecule and the method further comprises reverse transcribing an annealed RNA molecule into a cDNA polynucleotide.

16. A method for detecting the polypeptide of claim 10 in a sample, comprising:

a) contacting the sample with a compound that binds to and forms a complex with the polypeptide under conditions and for a period sufficient to form the complex; and

b) detecting formation of the complex, so that if a complex formation is detected, the polypeptide of claim 10 is detected.

17. A method for identifying a compound that binds to the polypeptide of claim 10, comprising:

a) contacting the compound with the polypeptide of claim 10 under conditions sufficient to form a polypeptide/compound complex; and

b) detecting the complex, so that if the polypeptide/compound complex is detected, a compound that binds to the polypeptide of claim 10 is identified.

18. A method for identifying a compound that binds to the polypeptide of claim 10, comprising:

a) contacting the compound with the polypeptide of claim 10, in a cell, under conditions sufficient to form a polypeptide/compound complex, wherein the complex drives expression of a reporter gene sequence in the cell; and

b) detecting the complex by detecting reporter gene sequence expression, so that if the polypeptide/compound complex is detected, a compound that binds to the polypeptide of claim 10 is identified.

19. A method of producing the polypeptide of claim 10, comprising,

a) culturing a host cell comprising a polynucleotide sequence selected from the group consisting of a polynucleotide sequence of SEQ ID NO: 1-1009, a mature protein coding portion of SEQ ID NO: 1-1009, an active domain of SEQ ID NO: 1-1009, complementary sequences thereof and a polynucleotide sequence hybridizing under stringent conditions to SEQ ID NO: 1-1009, under conditions sufficient to express the polypeptide in said cell; and

b) isolating the polypeptide from the cell culture or cells of step (a).

20. An isolated polypeptide comprising an amino acid sequence selected from the group consisting of SEQ ID NO: 1010-2018, the mature protein portion thereof, or the active domain thereof.

21. The polypeptide of claim 20 wherein the polypeptide is provided on a polypeptide array.

22. A collection of polynucleotides, wherein the collection comprises the sequence information of at least one of SEQ ID NO: 1-1009.

23. The collection of claim 22, wherein the collection is provided on a nucleic acid array.

24. The collection of claim 23, wherein the array detects full-matches to any one of the polynucleotides in the collection.

25. The collection of claim 23, wherein the array detects mismatches to any one of the polynucleotides in the collection.

26. The collection of claim 22, wherein the collection is provided in a computer-readable format.

27. A method of treatment comprising administering to a mammalian subject in need thereof a therapeutic amount of a composition comprising a polypeptide of claim 10 or 20 and a pharmaceutically acceptable carrier.

28. A method of treatment comprising administering to a mammalian subject in need thereof a therapeutic amount of a composition comprising an antibody that specifically binds to a polypeptide of claim 10 or 20 and a pharmaceutically acceptable carrier.

SEQUENCE LISTING

<110> Hyseq, Inc.
Tang et al.

<120> Novel Nucleic Acids and Polypeptides

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<150> 09/491,404
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<212> DNA
<213> Homo sapiens

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<212> DNA
<213> Homo sapiens

<400> 18

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<212> DNA
<213> Homo sapiens

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<213> Homo sapiens

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<213> Homo sapiens

<400> 35
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<210> 36
<211> 1392
<212> DNA
<213> Homo sapiens

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<210> 37
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<212> DNA
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<211> 1511
<212> DNA
<213> Homo sapiens

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<211> 2672
<212> DNA
<213> Homo sapiens

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<210> 40
<211> 717
<212> DNA
<213> Homo sapiens

<400> 40

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ttcctctccc tgctcgaacc tgctgtcagg gaaatccacg cccaggagga ccgtctcatc    360
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<210> 41
<211> 1424
<212> DNA
<213> Homo sapiens

<400> 41

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gaccgtggag tgttggtaaa tctaccatga taaactacct ccttgggctg gaaaatactc    360
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<210> 42
<211> 766
<212> DNA
<213> Homo sapiens

<220>
<221> misc_feature
<222> (1)...(766)
<223> n = a,t,c or g

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<210> 43
<211> 849
<212> DNA
<213> Homo sapiens

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<210> 44
<211> 1476
<212> DNA
<213> Homo sapiens

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<211> 1712
<212> DNA
<213> Homo sapiens

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<223> n = a,t,c or g

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<210> 46
<211> 755
<212> DNA
<213> Homo sapiens

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<210> 47
<211> 2820
<212> DNA
<213> Homo sapiens

<400> 47
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<210> 48
<211> 1517
<212> DNA
<213> Homo sapiens

<400> 48

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<210> 49
<211> 1614
<212> DNA
<213> Homo sapiens

<400> 49

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<211> 659
<212> DNA
<213> Homo sapiens

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<210> 51
<211> 450
<212> DNA
<213> Homo sapiens

<400> 51
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<210> 52
<211> 1044
<212> DNA
<213> Homo sapiens

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<211> 1328
<212> DNA
<213> Homo sapiens

<220>
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<222> (1)...(1328)
<223> n = a,t,c or g

<400> 53

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ttaatagtta gaagtaagaa ctactgctta taatatctgg gcagtgatca accatttcag    180
caactggctt gctactaccc tcagcatatt ttatttgctc aagattgcca atttctccaa    240
ctttattttt cttcacttaa agaggagagt taagagtgtc attctggtga tgttgttggg    300
gcctttgcta tttttggctt gtcatctttt tgtgataaac atgaatgaga ttgtgcggac    360
aaaagaattt gaaggaaaca tgacttggaa gatcaaactg aggagtgcaa tgtacctttc    420
aaatacaaca gtaaccatcc tagcaaactt agttcccttc actctgaccc tgatatcttt    480
tctgctgtta atctgttctc tgtgtaaaca tctcaaaaag atgcagctcc atggcaaagg    540
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tctgttatgt gccatttact ttctgtccat gatcatatca gtttgtaatt ttgggaggct    660
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attgtgtgtc ttctagcaga aaacaaactg gtggtgtatg aaacatttta tatttcttac    900
tgggttttct gtaatatatg tatatgaata atttccacat gtatacctag aaaagtcttt    960
tacctaaagt tagtctacaa aagtacatat atatagatgg ctgtggtgtg accgtgtgtg   1020
cacatatgtg aatgtgtata tatcacgcaa caggagtgtc attcatgctg ctggcccctg   1080
gtgaagtgac aagtacaatt aaaggtggct ctgatccttt taaacaccta ccaaacccta   1140
aatttgattc caaaaggacc attctgcaaa gagtttgcaa agatctgggc ccacttgtga   1200
gcaccaacct ttaaacatga tgcgccagtc tcccaggagg ccctactcat tcccctacat   1260
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acagaagc                                                            1328
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<210> 54
<211> 804
<212> DNA
<213> Homo sapiens

<400> 54
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gccgagggcc agcgcagtgc cacgtcacag gccatgcacc agctcttcgg gctgtttgtc 180
acactgatgt ttgcctctgt gggcgggggc cttggaggca tcatattggt cttatgcctc 240
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gcctcgcaga tcctccccta ccaccaccag ggctcctgct gaagctaccc tttctggact 360
ccccccccaga ctcccagcgc tacgaggacc aagttcactg gcaggtgcct ggcgagcatg 420
aggataaagc ccagagacct ctgagggtgg aggaggcaga cactcaggcc taacccactg 480
ccagcccctg agaggacacg ctcctttcg aagatgctga ctggctgcct actaggaagt 540
tctttttgag ctccccattc cctcccagct gcaagaaggg agcccatgag cccagaagga 600
ggcccctttc cacaggcagc gtctccacag ggagaggggc aacaggaggc tgggaaatgg 660
tggggagtgg ggccgtaact gggtaccata gggggaaacc tcaacaaatg cccaacccga 720
ctgggcctaa ccagcctgca catggggtaa aaaaaggcca aattgagggc acccaagtga 780
atccactggc ccccacgtca acat 804

<210> 55
<211> 532
<212> DNA
<213> Homo sapiens

<400> 55
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gggagttacg ccccgctggt tttcggtgta ctccccgggc tggtgggcgt cggaggtgtg 180
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caggcagtct tgctgtcccc aggaggccct ggctccactc tggacaccac ttgttgatta 480
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<210> 56
<211> 957
<212> DNA
<213> Homo sapiens

<400> 56
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tccctctcct ccctccttta gttttttttgc tcttgtctca tctgctcagt gaggtccccc 180
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aactgtctct gaagaaataa gaaaaatagt ggtttgactt ttacgagaca actatgttta 840
ttattttgcc cttgcacatt aaatggctaa atttggccaa gcccctatct ccagaatttt 900
ccaggtaccc ctcatgttta tgtgcacagc aaaaggaggg cctttgctca tacttcg 957

<210> 57
<211> 410
<212> DNA
<213> Homo sapiens

<400> 57
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gtcggcaggc tcctggccct ggccttctcc agccccgcag ctccttgtgt tcacaccagc 180
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cgcaggctcc ttcccagtgc cctcagtccg ggaagctcag acagggagct ccaggaaatc 360
ctctaaaagg ggcccctggg aatactggcc acaaggtgga ggctctgccc 410

<210> 58
<211> 871
<212> DNA
<213> Homo sapiens

<220>
<221> misc_feature
<222> (1)...(871)
<223> n = a,t,c or g

<400> 58
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caaagacata aggcatttct atttgaggca ttggagaaat gagaggaatt gcatttgcca 180
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cgatttgggg ggaagggaac tagtctggga accttttggc taaattttag tcacttttta 540
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cctcttaagg acccagttta aagcccggcg c 871

<210> 59
<211> 636
<212> DNA
<213> Homo sapiens

<400> 59
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agctcagaag cccaggttga gagacccaaa ttgtca 636

<210> 60
<211> 996
<212> DNA
<213> Homo sapiens

<400> 60
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gaggaaacta aggaaactta attgatgact gaatgcagtg tgctgtgttg aactggatcc 960
tagagaaaat agacattagt ggaaaaacta ctgaaa 996

<210> 61
<211> 1622
<212> DNA
<213> Homo sapiens

<400> 61
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gcctgcagca gcagtgtgac ttgctcaagg tccagtgagt gacctcattc cccagaaaag 780
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ag 1622

<210> 62
<211> 887
<212> DNA
<213> Homo sapiens

<400> 62
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<210> 63
<211> 857
<212> DNA
<213> Homo sapiens

<400> 63
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aagagacctc atgacagctg gagttctctg gggactgttt ggtgtccttg gtttcactgg 240
tgttgctttg ctgttgtatg ccttgttcca caagatatca ggagaaagtt ctgccactaa 300
tgaacccaga ggggcttcca ggccaaatcc tcaagagttc acctattcaa gcccaacccc 360

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ttattctcag gtctggagca tgcagcagcc agaaagctca gcaaacatca ggacacttct     480
ggagaacaag gactcccaag tcatctactc ttctgtgaag aaatcataac acttggagga     540
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tgaaaatgct tgaggcttat cacctgccac agccagaacg tgcctcagga ggcacctcct     660
gtcatttttg tcctgatgat gtttcttctc caatatcttc ttttacctat caatattcat     720
tgaactgctg ctacatccag acactgtgca aataaattat ttctgctacc ttctcttaag     780
caatcagtgt gtaaagattt gagggaagaa tgaataagag ataccagggc tcaccttcat     840
ctactgcgaa gggaggt                                                    857
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<210> 64
<211> 2093
<212> DNA
<213> Homo sapiens

<400> 64

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ggatttgcta agaatatcag aaaaggagtg ttccaatttg aagagtatta caattgaaat    2040
aacatcaaat atgtcacact aagcagccag taacagaata aataattaca acg           2093
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<210> 65
<211> 683

<212> DNA
<213> Homo sapiens

<400> 65
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actggccctc cagaaaaaaa gag 683

<210> 66
<211> 1273
<212> DNA
<213> Homo sapiens

<400> 66
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<211> 2549
<212> DNA
<213> Homo sapiens

<400> 67

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<211> 533
<212> DNA
<213> Homo sapiens

<400> 68

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     <213> Homo sapiens


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     <211> 859
     <212> DNA
     <213> Homo sapiens

     <220>
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     <222> (1)...(859)
     <223> n = a,t,c or g


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<210> 71
<211> 864
<212> DNA
<213> Homo sapiens

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<210> 72
<211> 746
<212> DNA
<213> Homo sapiens

<400> 72
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<210> 73
<211> 1928
<212> DNA
<213> Homo sapiens

<220>
<221> misc_feature
<222> (1)...(1928)
<223> n = a,t,c or g

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<210> 74
<211> 3644
<212> DNA
<213> Homo sapiens

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<213> Homo sapiens

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<212> DNA
<213> Homo sapiens

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<211> 1929
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<212> DNA
<213> Homo sapiens

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<211> 654
<212> DNA
<213> Homo sapiens

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<212> DNA
<213> Homo sapiens

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<212> DNA
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<212> DNA
<213> Homo sapiens

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<211> 1301
<212> DNA
<213> Homo sapiens

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<212> DNA
<213> Homo sapiens

<400> 100

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<210> 101
<211> 529
<212> DNA
<213> Homo sapiens

<400> 101

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<211> 697
<212> DNA
<213> Homo sapiens

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<223> n = a,t,c or g

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<210> 103
<211> 711
<212> DNA
<213> Homo sapiens

<400> 103
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<210> 104
<211> 429
<212> DNA
<213> Homo sapiens

<400> 104
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<210> 105
<211> 1028
<212> DNA
<213> Homo sapiens

<400> 105
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<210> 106
<211> 738
<212> DNA
<213> Homo sapiens

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<222> (1)...(738)
<223> n = a,t,c or g

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<210> 107
<211> 1706
<212> DNA
<213> Homo sapiens

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<210> 108
<211> 851
<212> DNA
<213> Homo sapiens

<400> 108
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<210> 109
<211> 959
<212> DNA
<213> Homo sapiens

<400> 109
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<211> 435
<212> DNA
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<400> 110
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<210> 111
<211> 3545
<212> DNA
<213> Homo sapiens

<400> 111
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<212> DNA
<213> Homo sapiens

<400> 112

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<400> 113

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<212> DNA
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<212> DNA
<213> Homo sapiens

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<211> 1528
<212> DNA
<213> Homo sapiens

<400> 116
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<211> 1700
<212> DNA
<213> Homo sapiens

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<211> 445
<212> DNA
<213> Homo sapiens

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<211> 455
<212> DNA
<213> Homo sapiens

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<212> DNA
<213> Homo sapiens

<400> 121
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<210> 122
<211> 5186
<212> DNA
<213> Homo sapiens

<400> 122

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<210> 130
<211> 1245
<212> DNA
<213> Homo sapiens

<400> 130
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<210> 131
<211> 694
<212> DNA
<213> Homo sapiens

<400> 131
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gggccccccc ctctgttgag aacccaccca tgtc 694

<210> 132
<211> 466
<212> DNA
<213> Homo sapiens

<400> 132
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<210> 133
<211> 1845
<212> DNA
<213> Homo sapiens

<400> 133
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<210> 134
<211> 1019
<212> DNA
<213> Homo sapiens

<220>
<221> misc_feature
<222> (1)...(1019)
<223> n = a,t,c or g

<400> 134
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gggttcttat ccctgatgca catggcccct gctgctgtgt cattccccta ttggctaggg 180
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<210> 135
<211> 764
<212> DNA
<213> Homo sapiens

<220>
<221> misc_feature
<222> (1)...(764)
<223> n = a,t,c or g

<400> 135
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tattcactta agaaacaaat atatttagcc aggtacggtg gctcacgcct gtaatcccag 660
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<210> 136
<211> 1016
<212> DNA
<213> Homo sapiens

<220>
<221> misc_feature
<222> (1)...(1016)
<223> n = a,t,c or g

<400> 136
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gacaatgaag aatatttatt gagggtttat tgagtgcagg gagaagggtc ttgatgcctt 180
ggggtgggaa gagagaaccc ctcccctggg attctggaag tctaagtttc ccgtggtggg 240
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tcaaagtggc tgttgttcca gagttggagc catagaatcg tttatggatc cctgaaggcc 780

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acacggacac tgagggtggc tgagtcagct cataggaggc cacggatcct gtgcagtaag 960
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<210> 137
<211> 727
<212> DNA
<213> Homo sapiens

<400> 137
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gcgaggagga gtctgatctg catttcattt tgtcatctct gtgttctctc cattgggctg 120
cgtgtgattg tgtgcgttgt tgggatatct gaagatcgta aacgaagtgc cagtgcaccc 180
accctaggta ttgtacccct gcatgccagc cttcaccagc actgtgctcc aaaccaatct 240
aatccctgct cttggcatct gtgatctcta gaaagcgatc tgacagcaat cagaaaatgt 300
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tatagggagg tttatagaag acattgaata atataggttt aggcttactt ctcttagggg 660
aacatttttc tgacgtttat tactttgaag aggaaaaata tttaggatga cgaagctctt 720
tcttttt 727

<210> 138
<211> 659
<212> DNA
<213> Homo sapiens

<400> 138
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aacatgaacc cctctgccag cctcgtctgc ctcctctttg cgttttcttc ctgccgcatt 180
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<210> 139
<211> 2068
<212> DNA
<213> Homo sapiens

<400> 139
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<210> 140
<211> 580
<212> DNA
<213> Homo sapiens

<220>
<221> misc_feature
<222> (1)...(580)
<223> n = a,t,c or g


<400> 140
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ttgttcacat tgctgagttc agtcttttta gtttgcattt ggatatttaa gaccaatatc    180
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cctgtttcat ttatgatagg tagaataatc ccccccaacc ccaccaagaa atctggatcc    420
taatccctga acctatgact gggtggggca gcatggcaaa gggaaattaa ggttgcagat    480
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<210> 141
<211> 1276
<212> DNA
<213> Homo sapiens

<400> 141
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gcaaccacct tgtatttaca aaaggatcat gaagattttt ttaaacgaac attttcatag 180
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gtcaatttgc caatttctgt ggtgtaaaca cactcaccgc tgacacttga tagatgtttt 360
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<210> 149
<211> 1556
<212> DNA
<213> Homo sapiens

<400> 149
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<210> 150
<211> 688
<212> DNA
<213> Homo sapiens

<400> 150
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agatgctgga tcctccaggc tggggtagaa ttgcaacagc ttgtccttcc ttgtgggtgc 180
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cacacctgcc caagcggacg cgtgggtc 688

<210> 151
<211> 1667
<212> DNA
<213> Homo sapiens

<400> 151
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gttcaccagc tttcgggcta gctgggtagg aggtgatgct gccccggtct ggcacccact 240
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<210> 152
<211> 1040
<212> DNA
<213> Homo sapiens

<400> 152
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cgtagcggaa tcgggggtat gctgttcgaa ttcataagaa cagggaggtt agaagtaggg 180
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tatgcgatta ccgggctctg 1040

<210> 153
<211> 849
<212> DNA
<213> Homo sapiens

<220>
<221> misc_feature
<222> (1)...(849)
<223> n = a,t,c or g

<400> 153
tgaattagta ttgtactgca ttggaggctt atatagaaag cctttcccct agaaactggg 60
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ctatctgctg tgcctcagct gcttatttgg gacaggtatg gttacttata tatgcctggc 180
gtgctgaaac atctcttgaa actgagttct ataccattcc tttgtcttgg ctttactact 240
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ccagagcatt ttaaattaat cctttctgtt tcattattcc tcacttacac ttaaaatgac 360
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<210> 154
<211> 860
<212> DNA
<213> Homo sapiens

<220>
<221> misc_feature
<222> (1)...(860)
<223> n = a,t,c or g

<400> 154
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atcccatttg tgggttctct taattctatc attgcttctt ttcctgcgga aaagttttaa    180
gttttatgca gtctcatttg tgtgttttgc ttttgttgcc ttttggaata atctacagaa    240
aatcatagct caggccaatg tcatacagtc tccttctata tttccttgta gtagttctac    300
atttaaactt taattttgat ttgatgcttg tataaagagc aaaataaaag tcaaatttta    360
ttcttctgtg cccaaaaaca ttattgaaca agaccaagaa cacttaaaac ggaaacaaat    420
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cccaacgcgg agaacccccc tctctactag nnnacccaaa aaanannnaa ggggcgcccc    600
ttctggagga tcaaacttta cccgcccgcc acaaccaaac cttatccctt tcctaacggc    660
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ccccaattaa tccctctgaa cacgcccacc cgaaacaccg gacccgcgca acggacccgc    780
cgccctcacc acacgaaccg cctccgaccc ccccgcacac tgcaccgccc caactgccag    840
cgccgaagcg caccgccccc                                                860
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<210> 155
<211> 552
<212> DNA
<213> Homo sapiens

<400> 155
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cgcgtccggg ctgcagcacc cagggaggaa cgccgcggcc ctgttttttt atcatgccag     60
gaggctgcag caccagggaa tctgtgctca cgtcttccag gacagtgctt cttctagaag    120
ctgacatgga gctgaccaca gctcttggag gcatggcctg aggcttagaa aatagacaga    180
gatcatctga gatttcagca gtggggccac gtggcagcgc ccgaaggcct ggagcaggag    240
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gacctaaggt tgctcagtgg gccatggaaa cggcagtgat tggggtggtg gtggtgctgt    420
tcgtggtgac tgtggccatc acctgcgtcc tctgctgctt cagctgtgac tcaagggccc    480
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ctctcttcac gg                                                        552
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<210> 156
<211> 1120
<212> DNA
<213> Homo sapiens

<220>
<221> misc_feature

<222> (1)...(1120)
<223> n = a,t,c or g

<400> 156
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tgaaatgcag tgattaaagg acacaaggcc tcagtgtgca tcattctcat tgtggctttc    120
aggcggctgt ggaagacagg gtggggatgg tggcttcggg aggtgaggtg ctctgggact    180
tgggcaagtc ttaagcaagc cattcctgct ttctgggcct ggctcccatg ggccattaga    240
aatgaaaatg ctttgtggac tgctgaggac ggtgcaaggg gtgaggtttc cccagctcac    300
ccggatccat gggcccagca cccaggggca tcagcttctg cttttatggg tgggggtctt    360
gcaggttggg aantcgtcct tggccttca gaatgacctc atggggccct ccctgggaag     420
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cctttcgaag acttggcagc cgagcaccca cgggattgca tcagctccgt gatggctaag    540
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caggtccagg tacagctcca ggcccatagt ggggaccgac cgacaaattc cncgncnctg   1080
gcctaaggtc tcgatggnnn tccattnnnn ccggggggcg                         1120
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<210> 157
<211> 392
<212> DNA
<213> Homo sapiens

<400> 157
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gcaaaaggat gtacactctc accacttcta tttaaccttg gactaaaagt tccagccagt    120
gcaataaggt aagaaaataa aaatacaaaa atcaacatac aaccaactgc aaaggaaatt    180
ttaaaaaatt acattcacaa atagcataaa aagaataaag gatttagaaa taaagttaat    240
gaaagaagta caggacagta cactgaaaat tataaaacat tgtcaaagga aattaagacc    300
taaataaatg gagatatgtc ccatgtttgc aaataggaaa atacagtatc atcaaggtgt    360
cagttttccc aaaattgatc catagattca at                                  392
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<210> 158
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<212> DNA
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tctcccatcc tctcctcccg ggggctcaga gtccagagga gagtgaatct tgctgactga 480
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ggtccatgtt gcccttcatc attaagtcaa attccaaagc cccgggaggt tgtgaaggtt 600
cactcgcccc tgacgggaac gagacccagg gacttctgcc ccaccaggca tcctcggtgt 660
gggttgtatt tagagatggg cctggacagg ggccactttg ggcagccttg gttgcaagtc 720
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tgatggtgcc ccggcgc 797

<210> 185
<211> 1735
<212> DNA
<213> Homo sapiens

<400> 185
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tttggaaggt actggcttgg acggctgctg ggtgagtcct ttggagtgat gatgtcatga 180
tgtgggaaac gggccttatg gcttgtggaa acagatgccc tgtgttctga ccaaacaagg 240
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cattctgctg tgcagaccac cttttaaaag tgatcacaaa ccatttgctg aatacttgtg 360
gaacttgaat cctcaccaat gtctccattt tctggaatcc atcccaaccc ~~ccaccttggt~~ 420
cttttggaaa attgggctgt ttgctctttt tttcccctcc tctctgactt cttggatatg 480
cattgatgtt ttccccttcc ttccaaggaa ttataaccaa agtaaggtgt gtgtgtgtct 540
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gaatgcgggc tgggcgcggt ggctcacgcc tgtaatccca gcactttggg aggctgaggc 660
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gcccaggtgc ggggggctca tgtctggaaa tccccatcat ggtgggaggg cccagcgtgg 1680
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<210> 186
<211> 669
<212> DNA
<213> Homo sapiens

<220>
<221> misc_feature
<222> (1)...(669)
<223> n = a,t,c or g

<400> 186
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cccctgagcc tggtccttct cgtgtgtggc tggatctgcg gcctgctcag ctccctggcc 180
cagagcgtgt ctctgctgct tttcaccggc tgctacttcc tgctggggag tgtcctgaca 240
ctggcggggg tcagcatcta catcagctac tcgcacctgg cctttgcgga gacggtgcag 300
cagtatggcc cgcagcacat gcagggcgtc cgcgtcagct tcggctgggc catgggcctg 360
gcctggggct cctgtgcctt ggaggcattc agcggaaccc tcctgctctc agctgcctgg 420
accctcagcc tgagcccccc aatctgtggt catctgagtc cccagcaggt gggagggaga 480
gggggagact gaggcccaga gcggcagagg gacccaccca gatcgcctgg cgccagagag 540
atgccgtctc aggccaaggc ctccctggcc tctgttctgt ccactctccc cgaagggcag 600
gcttggtgga gaagaggctg atgagagggc ccgagagccc cttcgatttg cannnnnnnn 660
nnncaaggg 669

<210> 187
<211> 1804
<212> DNA
<213> Homo sapiens

<400> 187
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aaggaaagag ggaaccccag acgctggccg tccaaaatcc accaaagaaa gtgacctctg 180
agaaagtgag ccagaaacat gctgagcctt tgacagacac tggctctgag accccgactg 240
cccgacggct ctacactgcc agcgggcctc ctgagggcta cgtcccctgt tggccggagc 300
ccagcagctg tgggagcccc gagaacgcct ccagcgggga tgacacagaa gatcaggatc 360
ctcatgacca gccaaagaga agaagaatta ggaagcataa atcaaagaaa aaatttaaaa 420
atcccaataa tgttcttata gaacaagcag aattagagaa acagcagagt ctgttacagg 480
agaaatctca gcgacagcac acagatggca ccacaataag caaaaataaa aaaaggaaac 540
tgaaaaagaa acagcaaatt aaaaggaaga aagcagccgg cttggcagca aaggctgctg 600
gtgtcagttt catgtaccag cccgaggaca gcagcaatga aggggaaggc gtgggagagg 660
cttgtgagga ggatggtgtg gacaccagcg aggaagaccc gacactggcc ggggaggaag 720

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ccaggcagga agagggtgcg gacgctagtg aggaagatcc gacaccggcc ggggaggaag    840
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ccggggagga agacggtaaa gacaccaggg aggaggacgg tgcggacgcc agcgaggaag    960
acccgacatg ggctggggag gaagagggtg cagactccgg ggaggaggac ggtgcagacg   1020
ccagcgagga agatgataca attaccaatg aaaaggcaca cagtattcta aatttttga    1080
agtcaacaca ggaaatgtat tttatgacg gtgtctccag agatgcagct tcagctgccc   1140
tcgcagatgc cgctgaggag ctgctggacc gcctcgcgtc acacagcatg ctgccctcag   1200
acgtgtccat cctgtaccac atgaaaacgc tgctgctcct gcaagatact gagagattga   1260
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tctcagcttt ctttagttac tggatcacac atatccttcc tgagaagagc agtgactaaa   1380
atggaatatc tctttaagaa cagctcctct ttaacaaaa aacttaaaag acaaatgtga   1440
gatgggctta gagttagttc tctgggaact tgaaagacat ttatgccata ttatttattc   1500
acgtgtttgt tcctggtggg caagatgcca tctgaggctt cagatgagaa attggggtaa   1560
aatggaaatt tttcacttat ttgcaattat atatatcttg aattactaca taaaacttga   1620
ttctgtttct ctacttattg taaaaattga aaatggacat tctgttaagt taaatgtata   1680
gtttgaagct catatatttt tatgaagttt tgaatcacct tgtatctgaa agtctctgct   1740
ttaagaatgc tttctgggta ttaaaatgtt ctagtttaag tagtttgaaa aaaaaaaaa    1800
aggg                                                                1804
```

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<210> 188
<211> 1070
<212> DNA
<213> Homo sapiens

<400> 188
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tgtgactcag gattttaaaa atctggtgga cttatgccgt aagggagcat tttcctttaa    120
catttgtttc gacatagttt gccctggcgt tgttcagttt tttttggagt accactaatt    180
tctcccatac ctatgagcag gtagtatgaa ttttccattc tgggagagac tctattgtag    240
ctaaactgcc tgtattcaag gatgccttac ctcattttat tctttgctgt gtacatattg    300
tataagattc ttgtcaaagt ccatcttttc atagcagaaa ttgcccttta tgatttttta    360
aaattctttg agttatatgg aatctgcatg tttaaaacac ttacctgtct ggtagtgact    420
actctgatat ttattaatct acttagtttg taagtaaagt aaacatttac atctggttaa    480
aatttactat acccccccca aaaaaaaact acctgtttgt ttacctcata actgattctg    540
tttacatata cccacacata cacaacccac caatactatt aagcttttaa tgtggacatt    600
ccaataagaa aacagatcat tctcattgac tcttactttt tgagatgtat ggccaaattg    660
taatttatcc tggctacaaa aagaagaatc taggcaaaga ctaaagaaag ccaattgtca    720
tgacacagtt acactaggat tagactttgt taaaaaataa ctccacaagg atttgcaatg    780
gaatttcaaa cattatcttg gggaattctg gagaaaagac cattttactt agacctttat    840
gtttttgatg gtgctgtgca agagagaagc caggattttt tcagaaacac tcaaatactg    900
gccagacgca gtgggcgcat gcctgcaatc acaacactct gggaagccaa ggcagaaaga    960
tcgcttgagc ccaggagttt gagactagcc tgggcaacat agggagaccc cgtttcttat   1020
taaaaaaaaa cctgggggtt gggggccctg cctgtgggcc catttaataa              1070
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<210> 189
<211> 863
<212> DNA
<213> Homo sapiens

<400> 189
cggcccgtaa ttaccggctc gacgatttcg tcgctgacta gggacagggc tgtcacactg     60
ccccaggagg aatggaagct ttcccgccaa cctgcctcct tcctctggac tccctgtgtt    120
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ggtttatgta cttcaatgtg atacatcagc agtctctttg gtctgggctg accttccaca 180
ttggttggtc tgtctgcccc tcccttggga tggcgcttgg tgtcagagtg tggggaccac 240
ctccaggaca agcgccactg ttgtgcgcag ctcagccaca ctgctctggc ctcagtttcc 300
cctgtgcgga atggggatga gaatgcagtc gagggaggcg aggagctgca gtgctgaggg 360
ctgaggagtg agctgagggc ttaacccccg gcgccatcct tggagggagg gagggagcaa 420
tgcgaccggg gggccttggc taatcatcta accgcagatg tcacccccca cactgatatg 480
tgatcacgtc agctggccct gggacggtca gataccttgc acatgatgct gggtccgcca 540
gaggcaagac tctctctctg cattttactt tggatctcca tcctttgtcc atggtacagg 600
ttcaccctgt attgttcatc ctggccctat cctatctttg actcgggata ccgacccttg 660
tttggcacaa cactcctttt ttaaacctaa ctttctgtgc cggattccag tttaagcaac 720
cggaacctaa gctgaaaccg aaccacccta actgggggc caaagcccga actaataaac 780
cggttacggt accgcccctt gcgataatac aaaaaccgtt ttgtgctgcg ccctgaaaga 840
acgtgcccca gttaggcctt cac 863

<210> 190
<211> 420
<212> DNA
<213> Homo sapiens

<220>
<221> misc_feature
<222> (1)...(420)
<223> n = a,t,c or g

<400> 190
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actagctgta cccaccttgc ctggggcccc cgtgctgggg gtcgcccccca agatggtggc 180
ggccccaggg aggactgtac tgccagcccc agcctctggc cgctaggcac cccctgcctt 240
gccctggccc ctcactccga ggccagcgcc atgctgcgcc tggggctgtg cgcggcggca 300
ctgctgtgcg tgtgccggcc gggtgccgtg cgtgccgact gctggctcat tgagggcgac 360
aagggctacg tgtggctggc catctgcaac caaaaccagc ctgcctacga gaccatnccg 420

<210> 191
<211> 988
<212> DNA
<213> Homo sapiens

<400> 191
gctggcgatt tctacactgt tgcccgggct ggagtgcaat ggcacgattt ctgctcactg 60
caacctccgc ctttcatgtt cacacaattc tcctgcctcc tgagtagctg ggattacagg 120
cgcacaccac cacacctggc taattttttt gtatttttag tagagacagt ttcactatgt 180
tggccagact agtcttgaac tcctgacctc atgatccgcc tgcctcagcc tcccaaagtg 240
ctgggattac agttgtgagc caccgtgccc ggcctcagtt atttttaaag caaatctaga 300
tatgttttgt taagggattt ttaaatttc ctaaaaaaag ataacctgca atttcttgcc 360
ccaagtcatt ccctactgac aaattgccca tcttcctgat tctctgatcc cctcctttct 420
cctcattttc caaattcaga caagtctgtg catggggtga tatcaccgca ccagctctgt 480
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gaaaacgatc cttcttaatg accatgaatg ggctggagtt gcttggcaat cctgtcctct 600
gcaataggac atttaatctg cttgtggcct ttcgccatgg tggtggcgct cttcccttat 660
ttgggttatt tttctggatc cctttccact caaatcgggt cagaccttcc ctgacactcc 720
ttgtacactg caaattcaca ttaagcatat tatgtttcac atagtccaaa tgaaacagtg 780
atttggctac tcatttatta actgtccggg agttccagca gggtcacaat cacggctgtc 840

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tgtaccgtcc gtgagtgctc aacatcaccc agcacagggc ctggcactca gtaggtgctc     900
agtaaccatg cgctgaatga atgagtaaat gaagggaggg atggaatgaa ttgcaacccct    960
tgataactgg gacaattatt catggagg                                        988
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<210> 192
<211> 967
<212> DNA
<213> Homo sapiens

<400> 192

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gggtggaatt cggaaagtga tacaaaagat tactagccat actcattgca gatttcatga      60
agagagggtg agcatttgaa gcatttcagt ttgctattct ttgggggttg gagaatgcat     120
tccaatctac ctaaaagtgc cctttccctg gctgtttggg tgataacatt ttttgagctt     180
tggcagaggt tttaaactct gtatgtgggc tggatatgtg atctacacac tgttttgtag     240
gttttctttt tctctgattt caattagaat cagaaaactt ggcagtattg ggtttgaatt     300
gccacttggc aataatagtc agctgggttg ccccctttaa aatagataag cattctctag     360
tttgccacag gtgacactac cccccattgcc tcttcagctc actcattcac atttcctgat    420
gggcatctgc aggtgtatct ttgaccgctg tctggatgtt ggaatgagtg gttcgctgag     480
cagacagcct gactcctgtg tatctcccat gattgtccaa gcatcactta ttgctccttg     540
accctgtctt tttactgacg tagttgagtg ttgtgcagcc ttttatttta gaggcagggt     600
ctcgctctgt cacccaggct ggagtacagg cgcggcacaa tcacagctca ctgcagcctt     660
gaactcctgg gctcaagtga tcctcctgcc tcagcctccc aaggattata ggcgattgcc     720
accatgccct gctaattttt tatttttagt aaagataagg acttgctgtg ttgcccaggc     780
tggactctaa cccctgggct caagcagtct tctcaatgtg ggcatcccca aagcgttgcg     840
attatgggta tgagccattg cgccctgcaa gttggcatac ttctaaattt tttgggaggg     900
tcctgcccaa ggcagaaggg aaaattgggt tgtagggctt gatgtgccca ggggacgtta     960
agcgcct                                                               967
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<210> 193
<211> 2238
<212> DNA
<213> Homo sapiens

<400> 193

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tttttttttt ttgatgattt ggatattatt attacaaaga atttaaatat acaagtttgg      60
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agaagtatca taagagtcaa acagttaaac atttctctgt gctttttttt tctattttct     180
aggaaatgtt gggtttagag agaagctcat caacttactt atacaaatca ggatatactg     240
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gaaatgactg agaagataca gtgatatgat atttaaagca aatattggca tatgcttata     360
caagaaaggc atcttacaat aatatttctg ttggtacatt acaatttttc agctagtaat     420
tctaaaatgc cagaggtcct atgatgcaat atcaaaaaaa ccagggaact gacatacaaa     480
gtcaaatata aagatagtaa cattcagtca tccacagata aaaggctatc tggacataag     540
cctgaaacaa gcaagacgcc atccactgcg atttcgccgg ttttgccctt gccacgttct     600
gcttcaaaaa tgatgctttt ggtagcatca gttccttgat acaactgaat tttccctgtc     660
ttccactttt catcctcact cgtggtcttc tcccatgcca gggcattgtt actgtttttc     720
acaaacactc gaagtttccc gactttgtct ccggccagcc ggtaatcaaa gagcaaacag     780
aagttgcttt ggggttgcag gtcaggtagg agaagtttca atcggccaat gtctttcctt     840
gtgaccctgc caaaggccgg gaactgccaa tatagaagcc aaatagcatt atctccgatc     900
agcaggattc ccagtcaaaa tcatcttcgt ctatcctgtt tcccagtcac agatccccat     960
gattgaagct gcagtcaacc gagatattta aatctgcttt atgttccagt ttggaagtta    1020
gcgctttcct ttggaccgag aatcaggccg aattcacctg cttcattcac cttagggaaa    1080
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gatgccttgc actggccgcg ttcccgaaac cacctgccac ccaggagagc agcagcggga 2160
gcgcaaggct ccagggcaga ggcattctcg cacgggtcct ccttcctcct cctgagcccc 2220
cctcgggagg gcgccggc 2238
```

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<210> 194
<211> 3326
<212> DNA
<213> Homo sapiens

<220>
<221> misc_feature
<222> (1)...(3326)
<223> n = a,t,c or g

<400> 194
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gtaacagacg cttcccgggc tccaggcctc cgagtgcccc cccccgccca ctctctgggt 120
cggcgtacat tgggcccttt ttctctgtct ctcgatatct ctctggcctc aatcgtcctc 180
ttggcgagtc tctctgtcgt ttcagtctgt gtggatttca gtcaccgcct cactctgtca 240
ctcttcctgg tgctctctct ttttctttat ctgcagcata tctggaaatg cctctcccct 300
ctgtttattc ccagcccccct cctgcctgcc caccccttccc acagaaagaa tctcgagatg 360
gggaaactga ggctcggctc ggaaaggtga agtaatttgt ccaagatcac aaagctggtg 420
aacatcaagt tggtgctatg gcaaggctgg gaaactgcag cctgacttgg gctgccctga 480
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tcgtccacct gggggatccc atcacagcct cctgcatcat caagcagaac tgcagccatc 600
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tcacaaccag cagcctcatc tgccagtggg agccaggacc tgagacccac ctacccacca 900
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cacaactgtg tcttgatccc atggatgttg tgaaactgga gcccccccatg ctgcggacca 1140
tggaccccag ccctgaagcg gcccctcccc aggcaggctg cctacagctg tgctgggagc 1200
catggcagcc aggcctgcac ataaatcaga agtgtgagct gcgccacaag ccgcagcgtg 1260
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ctggccactg gagcgactgg agccccagcc tggagctgag aactaccgaa cgggccccca 1440
ctgtcagact ggacacatgg tggcggcaga ggcagctgga ccccaggaca gtgcagctgt 1500
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<212> DNA
<213> Homo sapiens

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<211> 845
<212> DNA
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<210> 222
<211> 1947
<212> DNA
<213> Homo sapiens

<400> 222
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ggccaagtga aaagtcagta cagattctta tttttactat taaaaaaaa aaatcaaagg 180
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<210> 223
<211> 1131
<212> DNA

<213> Homo sapiens

<220>
<221> misc_feature
<222> (1)...(1131)
<223> n = a,t,c or g

<400> 223
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<210> 224
<211> 975
<212> DNA
<213> Homo sapiens

<400> 224
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<210> 225

<211> 1601
<212> DNA
<213> Homo sapiens

<400> 225
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ttgaacccac tgtcaacccc agctctggta taagcaagat ggagctgctg gcctgcttta 1560
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<210> 226
<211> 974
<212> DNA
<213> Homo sapiens

<400> 226
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gtgtgagttt ctttgtaata aaatgtatca atatgttata gatgaggtag aaagttatat 900
ttatattcaa tatttacttc ttaaggctag cggaatatcc ttcctggttc tttaatgggt 960

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<210> 227
<211> 666
<212> DNA
<213> Homo sapiens

<400> 227
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cgaggtgcac gcgaggctct ccacagccca ggaaggcagc ctgtcaccct gctctccgag 480
ccaggggcca aggtgtgggg ggcacaggcc atcctcatcc tgccaggccc ccgctttcag 540
gagtggggtg gtgccaatgc tcccactcag aaccctggac tgcggggtcc cctgagcaga 600
gggaccagcc agttccccat agacagattg gtgctggaca ggggctgcct gggccccagg 660
cttggg 666

<210> 228
<211> 1918
<212> DNA
<213> Homo sapiens

<220>
<221> misc_feature
<222> (1)...(1918)
<223> n = a,t,c or g

<400> 228
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agaggctgag gccagagaat cgcttgaacc tgggaggtgg aggttgcatt gagccaagat 1860
cacgccactg cattnccagc ctggtgacag agcgagactc agtctcaaaa aaaaaaaa 1918

<210> 229
<211> 1593
<212> DNA
<213> Homo sapiens

<400> 229
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tcttgcagct gaagggaggc actccttggc ctccgcagct gatcacatga aggtggtgcc 180
aagtctcctg ctctccgtcc tcctggcaca ggtgtggctg gtacccggct tggcccccag 240
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<210> 230
<211> 1583
<212> DNA
<213> Homo sapiens

<400> 230
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<213> Homo sapiens

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<212> DNA
<213> Homo sapiens

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<223> n = a,t,c or g
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<211> 1600
<212> DNA
<213> Homo sapiens

<400> 276

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<211> 1293
<212> DNA
<213> Homo sapiens

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<211> 1479
<212> DNA
<213> Homo sapiens

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<210> 279
<211> 1790
<212> DNA
<213> Homo sapiens

<220>
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<222> (1)...(1790)
<223> n = a,t,c or g

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<210> 280
<211> 5612
<212> DNA
<213> Homo sapiens

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ggccatagtg cccatggagg tttggactta agagatattc attggcagct caaagacttc   3360
caccctggag accacactgc acacagtgac ttcctgggga tgtcatagcc aaagccaggc   3420
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attccgctcg tgccgaattc                                               3620
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<210> 295
<211> 627
<212> DNA
<213> Homo sapiens

<400> 295
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tttacattga attccatgat atagcatatt tttaaaaata tgaaaatgat gttcataata 180
accaactggt tgaattatta ttttttgctg ttctcaccct ccaaccctca aatacaatcg 240
atcctccatg aagtggcgcc actgtggttc agaacacttt acactttgct tagagggtgc 300
tccacctgga agggcctgag ctcctaaaca atcggtaatg cagtgataaa gcgttaactt 360
ccaactatca aaaagtacct gactcattca ttccaactgg agctcatccc cgtgagctct 420
gggtcagaga gatgagctcc ccagccctgc cacagcgtca tgccaggaac caaactaaca 480
cgagcctcag gctgctgatc ttaaagtggg gatagcctta gggtcatctc ggcctctggt 540
gagccatcat ggcagcctct cggcagggtc tgagtggcag gagagcctcg gagagcctta 600
gaactgcctc tgttcttact tggaaac 627

<210> 296
<211> 888
<212> DNA
<213> Homo sapiens

<400> 296
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cagtatttac tgcctctaga taatagttta ttaaatactc tcccagacta tataactgag 120
aaaatacact aacaaattcc cctccccctt ttctaaatta aaaacatagt atatatgaat 180
atcattttca tatatcttgc tacttcctta gccttcttaa ttataaactt gagtcagcta 240
ttatttactg agtacttaca ttttagatgc tgttctaagt gctccacatg tataaacttg 300
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taaagtttaa atactttctc caagttcaca tggctagtag gtgggggagt gacgatttaa 420
accccctgctc ttaatctctg tacttttctg tctgatgtaa atttcttatt gccctttttt 480
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gatattcttt atccagtttg tctaaagtct aaaaaacaaa acaactcttt ggtttattac 600
tgggtgaacc ccaaaattgg gattcggcca gagaggccac atgggttctc ggcttcctcc 660
aggaaagaat tcaagaacaa gctgacagta aagtgaaatc atgtttatta agaaagttaa 720
ggaataggcc cagcacggcc gactcacacc tgtaatccca gcactttggg aggccgaggc 780
gggcagatca ctgggtgagg agatcgagac catcctggcc ggcatggtta aaccccattt 840
taataaaaaa gccaaacatg gccggcgggg gggcggccct cggggccc 888

<210> 297
<211> 675
<212> DNA
<213> Homo sapiens

<400> 297
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ccctcctgat gggtttcctg atggtctgcc tggggccctt cttcatttcc tggggctcca 180
tattcgactg tcaggggagc ctgattgcgg cctatttgct tctgcctctg ggtttgtga 240
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cggctggggg tgacagccat ctcagaggac gcccagaggc gaggccaaga gtgctgaggc 600
agagaaaact tttccagcac tcatgatgcc accactgtgg ggagcagcta ctgttattaa 660

aggccaacga gggac 675

<210> 298
<211> 379
<212> DNA
<213> Homo sapiens

<400> 298
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cggcggaaaa ctcattctcc tggtgatcag cccatgacct acacctccag acaaaataaa 120
acggaaaatt tgctacaatc actaatgagg gatccatgtc cagtgggagt ccagcttcga 180
actacaaatg atggccataa aacctactat actcgtgaca cagggtttaa tactttgttg 240
gaaatgtcat aaaatgatat gctcttactt caacttacaa ctggaacgac actttctgga 300
aacaattcaa tccgattctt tcatggagaa acttacattg acagatttga cgatttacag 360
aattcatgtt gcgacccat 379

<210> 299
<211> 887
<212> DNA
<213> Homo sapiens

<400> 299
agtacccctc cgattttcgg tcgacccacg cgtccgcttt tctccccctg catttcctat 60
tatttccata tttggtctcc tggacttggc atccaggtct ctctactttt tcactcaaat 120
cattgaacct tagctccatg ccttgcagtg gttcttctgt tcagactttc agaccattac 180
tgatttttca taatgtgacc ttcttcattt tacctgttaa gtgttttaat gctctgatta 240
atgttttaga aagaccattc tggcagctgt tgggagagat tggagaggaa tacagaggaa 300
gtgaggactg gttaggaggc agtttcaggt gagagatatg gtggctcaga cagggtgaga 360
agatggagat gagagaacag gtaggatgga ggaatgcttt acatgcagta gccgtaggac 420
ttggcggtgg tttggacctg ggagttaaga gagtgggagg gggacaagga tgtctctcag 480
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tgaaagttag ccacttgggc cggatgcagt ggctcacgcc tgtaatccca gcactttggg 840
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<210> 300
<211> 935
<212> DNA
<213> Homo sapiens

<400> 300
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gtccttctga ataaaacata gttgtttata agtcttggtg tacctgactc actcatttta 180
gtgcatcgag gtaggtagat tggagggtga ctgaggggag ggcactgtca gttgtgaggt 240
tgtcttctaa cagagtatgt acaggaaggt aatagttgct ttaacagtgt tcagacttca 300
aaagtgtagc tgttggagaa gtaagagcat caagcaagga gtggaacact tttggttggg 360

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ttatgtggtt atgagattag agcattctcc tattggttgg tgtcttagtc agctcagggt 480
gccatacaaa ataccataga ctgggtagct taaacagcag aaatgtattt ctcacagttc 540
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cccagctcct ggtttgcaga ctgccacctt ctcagtgtgt tttcatgtag cagagagtga 660
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caggggggca atttgggttc atggaaccct tggcttccgg gttggaagga atttctggc 780
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aggggatccc cccacttaag cttcccaaag ggtgg 935

<210> 301
<211> 2283
<212> DNA
<213> Homo sapiens

<400> 301
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tgggaacagg tgaagcccct ccccacacat acactccggt ggatgtgagc gagggtcctg 180
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tcagcctttc tgccactctg gggtcagtga ggtcttccgg ggaagccaca ctcagccgca 480
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ccttccaggg tgagccaaga aggcagacca gcgtccagga ctcgcagagc tttctgaacc 2220
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acc 2283

<210> 302
<211> 413
<212> DNA
<213> Homo sapiens

<220>
<221> misc_feature
<222> (1)...(413)
<223> n = a,t,c or g

<400> 302
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acggtcgccg gtgcgaccta ctactaacga ggcagtatgt actgggtcac agtcatcacc 120
ctgatctatg gctactacgc atgggtaggc ttctggcctg agagtatccc ttatcaaaac 180
cttggtcccc tgggcccctt aactcagtac ttgatggacc accatcacac ccttctgtgc 240
aatgggtatt ggcttgcctg gctgattcat gtgggagagt ccttgcatgc catattattg 300
ggcgagcgta aaggcatcac aagtggccgg tctcaactac tgtggttact acagactttg 360
ttctttggga taacgactct caccatcttt gatgcttaca aacggaagcg ccn 413

<210> 303
<211> 681
<212> DNA
<213> Homo sapiens

<400> 303
cactggtgga attcgttctg aggagccaaa ggaggaagag actttcgggg aaagaggaga 60
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tgctgacttc cctttcctgg acttgagctg atgaagggga aatggtgttg cagtctcctc 180
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cctggtcccc agtgccctct cctctggctg gggcaggagg acctgccgga catagccaga 300
tgtattacgg atgactgcag tcagctcccc caggctcctg cttctcttgc ctcctgcttt 360
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gaacccagct ccctttgtat gactgaccct gccagcctgg gagacataga gtctgattgc 600
ccggtctggg ggttataacc ccccggggtt tggacctgga aatccaaagc accctttggg 660
gctaagacct gggccaagcc g 681

<210> 304
<211> 427
<212> DNA
<213> Homo sapiens

<400> 304
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ctggtggctc ttcatgctgg ctacatcttt atccagacgg agaagaccat ctacacccct 180
gattcactac cgggtgttca ctgtgaacca caagatggac cctgtgacca ggacattcac 240
tctggacatc aaggtggtct ttcccgatga ggggtggggg gtggtggtgg atcctggaca 300

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ctggggttac atggtgtgct gaagtcctgg gggcatgagc caccagggcc ctcccagagg    360
gcagtcacca gccccacccc ctatccccac agaacccaaa gggaaacacc gtgattagcc    420
agagtct                                                              427
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<210> 305
<211> 609
<212> DNA
<213> Homo sapiens

<220>
<221> misc_feature
<222> (1)...(609)
<223> n = a,t,c or g

<400> 305

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acagggtgtt tctggtgagc ccctaaacac cagcatggtg atatccactc agtatctttt     60
tacccatatg ggtgggaggc tttggatttt tctccagcta tgtcagagcc tgggtctgag    120
cacagtggtc agcagcagac ctgttgcctg tctggagtcc gttcctggga tgtgtatgtc    180
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tctttccaga gctcatcttt gaatccttgt tattataaaa taagaattaa attgttgaac    300
tataacactt agggttaacg ggcacataaa tacttttttt aaatttttta acatatatat    360
attttttca tcacatttc attgtattag gtatcagaat tttttttttt aattcagtac      420
agatttacgg cctggggggg gggctcacgc ttatagtccc aaagttctgg gattacaggc    480
gtgcacnctg tgcccggcct aacattaatt cttagttatg tgcacagtct tatgggcaca    540
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acaatgatg                                                            609
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<210> 306
<211> 608
<212> DNA
<213> Homo sapiens

<400> 306

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atcagctccg actggctttg cttgaatcaa gagtttgccc ctgttcaatc agccatagcc    180
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accagtgggg tgtaatcact agggacacct acatttgcct gtagtgtaga gagggactga    300
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aggtttggga tcctgaggcc agaccagtca ggcagtccac cctgaacaga gcccatgcag    420
gacagtgggc atgagacccc aaacctctgg ctgagaatat tgccctcact taaagaagga    480
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tctactaa                                                             608
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<210> 307
<211> 781
<212> DNA
<213> Homo sapiens

<220>

<221> misc_feature
<222> (1)...(781)
<223> n = a,t,c or g

<400> 307
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tctaccaaaa atacaaaaat tagccagcca tggcggnggg caactgtaat cccagctact 660
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<210> 308
<211> 1391
<212> DNA
<213> Homo sapiens

<220>
<221> misc_feature
<222> (1)...(1391)
<223> n = a,t,c or g

<400> 308
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acagaaagga atgtactttg caagaaatgg tagttcatcc taagtttcca aatacttttg 180
gaaggctaat gcagcagctg ggcaaaataa cacacagtac acaaagaaca gtgtatttca 240
cagagtcagt aatgaaaaac tgacagctct ttaggcagga tatgcttttt ttcatttttt 300
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cagtggggga aagcaaggga acaaattcca taccatcatc tgacactaat ggagtatggc 1200
attattaaaa aaaataaagc ttttgcattt taataacccc acagaaaagt ctatgagcaa 1260
aagacttgat ctgtttgcca ctcaaaagtt agagatctca cagtgaaatt agaaaactct 1320
aattatacat atttcggacg cgtgggtcgn ccctgcagat ggngatcatn ccgacgggat 1380
cagtgggggc c 1391

<210> 309
<211> 874
<212> DNA
<213> Homo sapiens

<400> 309
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<212> DNA
<213> Homo sapiens

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<212> DNA
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gggggctgga caggggggaag ggagagagga tttctattca cctttaatat atttttacaa  5820
aaaaagcaaa caatttaaaa acaagcccac cgcttctgta catgtctaaa tatattttta   5880
gaagtgggta ggattgtgaa tttctgatgc agggcctttt tataaatagg ttagggtagc   5940
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taatttatat ttttttttaga tcctcccttt cctatagaga taaaagtgat ttatcttggc  6060
aattgctttg cttggcattc tttttttttg tgatgagggt ggtggtgtgg tgcagggtct   6120
gggagtgctg ccttctcctt gtactctttg tctctccctc agcaagttgt caggcatttc   6180
cctggtgctc agccttatgc ttgaagtggg aagggtattc ccaccctcag gagggacacg   6240
cttcacac                                                            6248
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<210> 318
<211> 402
<212> DNA
<213> Homo sapiens

<220>
<221> misc_feature
<222> (1)...(402)
<223> n = a,t,c or g

<400> 318

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tggccgcctg ccctctctcg tgctggtggt gctgctggtg gtgatcgtcg tcctcgcctt    180
caactactgg agcatctcct cccgccacgt gctgcttgag gaggaggtgg ccgagctgca    240
gggccgtgtc cagcgcgccg aagtggccct ctggcgggtg ggagggcgca attgcgacct    300
cttgctggtg gtcgggacgc gcagtagacg gatcgaggag aggggagccg actacagccg    360
gctcagcagg cggctgcagn ccaaagaggg cctcgtgaat ag                       402
```

<210> 319
<211> 635
<212> DNA
<213> Homo sapiens

<400> 319
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tttcgtggag gctcagaaag acccctaagg agcgggtatt caatctagcc tcagaagatg     60
aaattcagta ggcgagaagt gttggaacca aaatcctcgt tctggagtca ttttatggaa    120
gcagctgctt tggcttgaaa tggcaagccc cgggacctct ccccacccag tgctttgatg    180
agggccaggc cagcatgtac tgccaccttc ccgtcctttc acctagccct ggacagtagc    240
taccttcctt gctgtaaagg aaaggccacg tttataccaa aatccagaat ctatctgcag    300
gaggcaaagg gaagtgggga gcccctggga tgaggatctg tgagggtggc tttccctgct    360
aagcagaaca tctgactgtc tcactcctgg ctgtgtccag gaggtagatg ggcttgaaat    420
caattctgct tgctgcatat ctgatttcct agagcccact cgtcaagtga ggagacatcg    480
tcagtgctgc agccggggat cgccatggag accataggac tggctgactc cgggcagggc    540
tccttcaccg gccaggggat cgccaggctg tcgcgcctca tcttcttgct gcgcaggtgg    600
gctgccaggc atgtgcacca ccaggacctt ttttt                               635
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<210> 320
<211> 1311
<212> DNA
<213> Homo sapiens

<400> 320
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ctatcagcca cataccacat agggaggcca cagatgggcc gtggtgggtg gaggtagcct     60
ttgcaccatg ttgagcagag acggctggct ctcctcaggg ctccggctgg aaggtgtata    120
ccggaaaggg ggcgctcgtg cccgcagcct gagactcctg gctgagttcc gtcggatgc     180
ccggtcggtg aagctccgac caggggagca ctttgtggag gatgtcactg acacactcaa    240
acgcttcttt cgtgagctcg atgaccctgt gacctctgca cggttgctgc ctcgctggag    300
ggaggctgct ggtattccta agatccctga gagccaaggc ccaaccagga tctctgcctt    360
cccccaccag aatccatggt ttggcagccc tccgccccat cacttcccac cctgggggat    420
catccagaga cttggctcag gggaggtgg gaagggggca gagacacatc catcctgcat     480
ttgtgcctaa aaatccctcc ctctgtacca gctgccactc tttcttcccg ggtcctcccc    540
aaccctcctc cattccatcc ccagagctgc cccagaagaa tcagcgcctg gagaaatata    600
aagatgtgat tggctgcctg ccgcgggtca cccgccgcac actggccacc ctcattgggc    660
atctctatcg ggtgcagaaa tgtgcggctc taaaccagat gtgcacgcgg aacttggctc    720
tgctgtttgc acccagcgtg ttccagacgg atgggcgagg ggagcacgag gtgcgagtgc    780
tgcaagagct cattgatggc tacatctctg tctttgatat cgattctgac caggtagctc    840
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cagcagctgg gatggacttg tgggtgactt ttgagattcg cgagcatggg gagctggagc   1080
ggccactgca tcccaaggaa aaggtcttag agcaggcttt acaatggtgc cagctcccag   1140
agccctgctc agcttccctg ctcttgaaaa aagtccccct ggcccaagct ggctgcctct   1200
tcacaggtat ccgacgtgag agcccacggg tggggctgtt tgcggtgttc gtgaggagcc   1260
acctcgcttg ttggggaagc cgcttccagg agaggttctt tcttgttgcg t             1311
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<210> 321
<211> 867
<212> DNA
<213> Homo sapiens

<400> 321
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cttccatggg accctgcagc tgggccaggc cctcaacggt gtgtacagga ccacggaggg 180
acggctgaca aaggccagga acagcctggg tctctatggc cgcacaatag aactcctggg 240
gcaggaggtc agccggggcc gggatgcagc ccaggaactt cgggcaagcc tgttggagac 300
tcagatggag gaggatattc tgcagctgca ggcagaggcc acagctgagg tgctggggga 360
ggtggcccag gcacagaagg tgctacggga cagcgtgcag cggctagaag tccagctgag 420
gagcgcctgg ctgggccctg cctaccgaga atttgaggtc ttaaaggctc acgctgacaa 480
gcagagccac atcctatggg ccctcacagg ccacgtgcag cggcagaggc gggagatggt 540
ggcacagcag catcggctgc gacagatcca ggagagactc cacacagcgg cgctcccagc 600
ctgaatctgc ctggatggaa ctgaggacca atcatgctgc aaggaacact tccacgcccc 660
gtgaggcccc tgtgcaggga ggagctgcct gttcactggg atcagccagg gcgccgggcc 720
ccacttttga gcacagagca gagacagacg caggcgggga caaaggcaga ggatgtagcc 780
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agcagagtca aggcatctca aaaaaaa 867

<210> 322
<211> 1144
<212> DNA
<213> Homo sapiens

<400> 322
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agagaaacac agagattcct tattggcaat ctttctgttc tcttatttaa agaaaaaagt 180
tgatttttct ccttaatctg aaacgtatgg ctgctctgta gagaaggttt gggagatgct 240
gaaatggggc gagaagggag cactcatcag ccttacacac ggctctgcta aggatcaggg 300
ctccaggccc ctcagcctcc tccccagcat ggcagcccct tccagcctct cctatcccca 360
ggcctgcagg ctaggatggc ccggccctca gccttcccca tcggggtctg tctgactctg 420
cccatggcct ggatctcccc gggtttagct gtgcccagct gtccccagta catacttcaa 480
gcccaaggct gcatcctaga catgaaaacc cgaggcagcc atggggagtc tgctgtgcca 540
ggggcccatg gctctcgtcc cttccaccct ctggctgagc ccaatcctcc ccgccaaaag 600
ttgacaccat gcacatgagg gacacggggt ggctccccaa agctgacggt cgacgcccct 660
gcagggccgt gatgccaagt cagggtctca gcaggccctg ggactcagtc cccacagagg 720
gcagggggtg acactcagcc ccggagaagg gcccctcaga gccctctgac agtgcccttt 780
cccggtgggc aacgctttct gccaggcatg cgctcccacc agattacagg aaggctgcag 840
gcagagtgtg cacaccggga tggcccctta tcccgcccag acaaaggcgc gcagggccct 900
gaggcagggc ccatgctgtg ctggagtggg tggagctggg aacagaaata cgtcctgcct 960
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cagcaacaac aataacagca ggctgggccc ggtggcttac acctgggatc ccagcacttt 1080
gggaagccga ggcaggaagg atcgcttgga ggcgagggaa ttaagaacag cctgggcaac 1140
ataa 1144

<210> 323
<211> 366
<212> DNA
<213> Homo sapiens

<220>
<221> misc_feature
<222> (1)...(366)
<223> n = a,t,c or g

<400> 323
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gggggaaaaa cagtttcttc ttgtttcccc gactatgacc ggacattata atacaattta 120
gccgaatggt cagacatcgt ggcatggatg accattattc tccagataga gacagtcatt 180
ttcttactct acctcgctcc agatacagtc agaccattga ccatcatcac agggatggca 240
gggattgtga agcagcagat agacagccat atcacagatc cagatcaaca gaacaacggc 300
ctctccttga gcggaccacc acccgctcca gatccacttg acggncttgt accaacctta 360
tggggt 366

<210> 324
<211> 839
<212> DNA
<213> Homo sapiens

<400> 324
cccacgcgtc cggcttttgg tgtgttggat aggcttttga gtagggagag atactatctt 60
gaattgtgct aataatttaa ctcaacagca tctaacaaag gcagtcttat tcttggatca 120
tgtgtacaga tcatagtctg aagtggaata agcagaatgt tgtcctcagt gtgagatgtt 180
atttagaaca cactggaaac attgtgatgt cattgtgcac tgaggcaggg aaatgttagt 240
ctacatttta tggaatatgt acttcaatgt ttgcattgta cctggagtga taaaaagcaa 300
aacaggtact caagacctgt ctgggctttg gcctttgggc acattccccc tcatcacctt 360
ccttcccact tggctgagct atggatgaga aaacctaggt caatagttca ccaactcacc 420
ttcaagccag gtgggctgac aagtcctcct ttgaccacag gaccccagcg cctgcatcca 480
gaagcatcta agatcctgga agtcaactta aattttcaat gaatgggcca gttgcagggg 540
ctcacacctg taatcccagc actttgggaa gctgaggcga caggattctt tgagccccgg 600
aatttgagac caacctgctt gggccaccta aacccatttc atcaatcaat cataatcgag 660
ggagggggcgg gattggagcc ctcattatta ggagctgagg ggggggccac tggaccccgg 720
ggtttgggtt gccgggcccc tattggcccg gaccctggga aaaaacgaaa accagcctcc 780
gcagaactcg ccaaaaaatg gggcgggcgt tgaaaacaaa ttttaacccg gcgggccat 839

<210> 325
<211> 677
<212> DNA
<213> Homo sapiens

<400> 325
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tgtttctttt tgagttcttc cttaccttag ttccaatgtg ggcatttcct ggagacaaaa 120
cttttgtttc acctgcatca tctttaagtt ttcttgatct gagttttctg cttttctgta 180
acagtgtatc tattggaaaa caataacaga aatctcataa tcctaaaatg ttaagcattt 240
tgctaatatt acacagagta tgtgaactaa cagaagggct agattttgtt tatcttgtac 300
atcttggaaa tctgtgacag cttggcttag attcagtttt agtgtactgt atttgaaatt 360
accgttatcc acaggaacag taactatagt ttgtcctaat ataacgaagt ctactttata 420
agttggctga gcatggtggc tcacagctgt aatctcagca ctttgggagg ccaacatggg 480
cacatcactt gaggtcagta gtttgagacc agcctggcca aaatggagaa accccatctc 540
aactaataat aaaaaaaatt agctgggcat ggtggcacac gtcctgtagt cccacctacc 600
tgggaggctg atgcaggaga atccattgaa cccgagaggt ggaggttgca gtgagccaag 660
atcgcaccac tccactc 677

<210> 326

<211> 517
<212> DNA
<213> Homo sapiens

<400> 326
tgcttggcac gaggcaggag gctgtctgga cacactgatt actcactcac cagcctccct 60
cttttgtcca ccagcccagc ctgactcctg gagattgtga atagctccat ccagcctgag 120
aaacaagccg ggtggctgag ccaggctgtg cacggagcgc ctgacgggcc caacaggccc 180
atgctgcatc cagagacctc ccctggccgg gggcatctcc tggctgtgct cctggccctc 240
cttggcaccg cctgggcaga ggtgtggcca ccccagctgc aggagcaggc tccgatggcc 300
ggagccctga acaggaagga gagtttcttg ctcctctccc tgcacaaccg cctgcgcagc 360
tgggtccagc cccctgcggc tgacatgcgg aggctggact ggagtgacag cctggcccag 420
ctggctcaag ccagggcagc cctctgtgga atcccaaccc cgagcctggc gtccggcctg 480
tggcgcaccc tgcaagtggg ctggaacatg cagctgc 517

<210> 327
<211> 992
<212> DNA
<213> Homo sapiens

<400> 327
ctggtcttga actcctgacc ttgtgatcca cccgtctcgg cctctcaaag tgctgggatt 60
acaggtgtga atcaccatgc ccggctagaa gagctttatg ttcatgatgt tgagatgaag 120
ttggggccag aagaagagtc agttgataaa agctaaagta tttttagatc ctgattaaag 180
aagaaggtaa tgggttgact tgagagagaa tgagcgttct gttatgggaa tgctcatatg 240
ggaaatgttc tgtctctttg tcaaaaactg caggaccacc tgttggtgac attggaggaa 300
ttcctgcttt gtgtgggagg gtgaactaga tgcctttaaa aaaaatttcc cccccacaga 360
cttgttttag atattttact gcttcagaga gggtcatgtt cacaccattc tccccttttg 420
taatttttca cacctccctg gctccccttt tataatttag aaagaggttt acaagtctgt 480
aactttttgt attagattta ctttgagaaa tcttgtactt aatttagtag gtcacagagg 540
gttgctgaat gactggaaac ttgtgtttct ttttccattaa gggctatttg ctgacttctg 600
aaatattgat gatttatttg actttagaat tttgcatact gaggggaaag catcttaatg 660
tatcatttaa agcaggagat actttcatac tatacctggg ttctcttggc tttgaagagg 720
agggtggtcc tgagatattg aaagattgca tgggtggcct gtcatcccca ccactttgga 780
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cgcggcaccc cctcctctgc taaaaatccg gaaatttgcc cggggcgggg gggggatgcc 900
ctatacatcc agtttctcct caggcgggcc cattatatta aaccctagcc ggccgctccc 960
tcgcccccgc gcaacaatat atctatccgc cc 992

<210> 328
<211> 894
<212> DNA
<213> Homo sapiens

<400> 328
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tgccactttt ccttttgatc agtgtcctcc tgccatcctg gcctccttgc tgtttctcaa 120
acatgccatg tatgttcttt cctctgcaca cctgtgcttt ttatgccttc agtgctcctc 180
cctagaggtc tacttgatct cttccctcac ttcattcaga tctgtgctga actgttaccc 240
accagagaga tcttccctga ccattcaata tcaaatatta ctccttctgt tacagtaggt 300
agctagtcag gcatgagcag ggcagaagag ggctccccte cctcaacaca caccaggaat 360
gacaggcaaa catcaggtga tggtcaggca gctgctaact gtttctctaa aatattaatt 420

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ggttgcagcc tgcaccaggg aaaggcagtc tccatatata cagaagcacc tgaagctggt    480
gatcagcagc ttcccatgag atctcaggaa ctgggtgagt gggctcaagc gtttgcacta    540
agaggcaaaa tgccagagtt tggtatgtga cctcctaagg acattcgact ggtaatggaa    600
gaacacctca agtgaacacg cgtacaactc cagtaaacac gttgcacatg gtccctttcc    660
caagtgctgg gaggctactg tgtgtgcaga cagcctgccc caagggaaga atcatgggag    720
atgggacacc aagatcctgg aagtatgcca acatataaaa ccccaagttg aaaggtcaaa    780
ccgtgcattt gtcttttcaa gttgcccact ttgccctctt ccaagtgtac cttccttccc    840
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<210> 329
<211> 423
<212> DNA
<213> Homo sapiens

<220>
<221> misc_feature
<222> (1)...(423)
<223> n = a,t,c or g

<400> 329

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gagctattcg atagtgaaga cccccggcag cgagagtacc ttaagaacat cctgcaccgg    120
ctttatggca ggatgctggg actccggccc tacattcaca aacagagcaa gcacattttc    180
ctccggatga tctatgaatt ctagcacttc aatgggggg ctgaactgct ggagaaccta    240
ggaagcatca tcaatggctt tgcgctgccc ctgaagacgg agcacaagca gttcctgggt    300
cgcgtgctga tccccctgca ctctgtcaag gcgctgtctg tcttccatgc ccagctggca    360
tactgtgtgg tgcaattcct ggagaaggat gccactctga cagagcacgt gatccggggg    420
ctn                                                                  423
```

<210> 330
<211> 18819
<212> DNA
<213> Homo sapiens

<400> 330

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gtaacttctg aagaactgaa tattataatt cagaatgtaa tgacctgggt tgtggctaca     60
gtgaccagta tattgtaccc agccatcaca aagtatgaaa aaagattgca aaataataca    120
tacccagtat ctgatgactc catcctctct tcagatagtt caagtttctg tagcacgtgc    180
agtgaagact ttacatatag aagctacaca tctgcaacaa ctaaaacatt tcaggcagaa    240
ccctgtgcat ttgtagttga cacgtcagta aggagaccaa ccacacctat aaaacctcct    300
cctgcacatg tggaaaaaac agttgtgggg aaaacatgtc acataaaagg acaatctata    360
atctctaaac ataaatataa taaaaccaac ttgctatatt catacсctaa gctcagaagt    420
tgtaaatcag atagtcacct tttagcatca tttgaaacag gcacaaaaaa atctaaggat    480
gctaccactg aaacagatag cttagggagt tcattgcatt gtgataaaac agcaaaagcc    540
atggatgaaa tgaagaattt aaaaaatgtt tttgttaact ttaaatgtta cttgaaaggg    600
gaaactgaag tgattttaga aagcattttg cgagaaataa tgtctgattt aacccaggcc    660
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gaaatcttgc tttccaatgc tcatattccc tcagttgctt ctgagattgt ggaaaatatg    780
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aaacctttga aaaattcaat gcctcatact ttggacccaa tgtgtgatat tgcagaggac    960
atggtgcatg ccattttaga aaagctaatg actcttgttt cttttaagca aaatgaattt   1020
cttcatctta aagacacaaa taagctttcc tgccagcaac ataagacaga cccaatatgt   1080
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atgttccttc aaagagctgg caaaaataaa tctagtcttg aatctgatga agctagttta 1140
attgtcaatg aagaagtaca aaatttaata tcaaatattt tttcccagtc ttctttggtt 1200
gcttatatag aggaagcaat caatgctata ctaggttata tacaaactga actaaataat 1260
gagagaatta ttgcatctga agaaaccgta gtactccttc agctacttga ggacatcctt 1320
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agtagtcctt ctgacaccaa agaaaagtac agactcactg gcactagatt atcaaatagt 1440
cctaggtctg gaagaccatt tccacctata aatgttccag gcatggttct ttattctgat 1500
gatgaaaatg aggaaataga caatattgta aaaaatgtgc ttgattcaac tttcaaagat 1560
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aactacattt ccaataccac taaaagttcc atttcatcat cagttcatca gatttcctta 2100
cataattctg acactgaaca catagtcaaa gaagcaccaa ataaataccc attaaaaaca 2160
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ttcaaatctt cattacgatc tcaacttagt aagtacacag ctaaaatagt aaacattgtt 2400
ttatgtgcta tccagaatga actggaactt cacaaggaaa acctaaatct tagggagatt 2460
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tctcttgtca cgagtattga tgatgacatt ttggcgagtc cattattaac ctgtatttat 2580
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<212> DNA
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<212> DNA
<213> Homo sapiens

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<213> Homo sapiens

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<211> 2527
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<213> Homo sapiens

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<211> 332
<212> DNA
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<212> DNA
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<212> DNA
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<211> 279
<212> DNA
<213> Homo sapiens

<400> 342
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<212> DNA
<213> Homo sapiens

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<211> 440
<212> DNA
<213> Homo sapiens

<400> 347

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<211> 687
<212> DNA
<213> Homo sapiens

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<212> DNA
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<212> DNA
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<212> DNA
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<211> 409
<212> DNA
<213> Homo sapiens

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<210> 355
<211> 1449
<212> DNA
<213> Homo sapiens

<400> 355
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<211> 403
<212> DNA
<213> Homo sapiens

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<211> 794
<212> DNA
<213> Homo sapiens

<220>
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<222> (1)...(794)
<223> n = a,t,c or g

<400> 357
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<212> DNA
<213> Homo sapiens

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<211> 652
<212> DNA
<213> Homo sapiens

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<212> DNA
<213> Homo sapiens

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gttttgaatt ccaggcaaaa aaagtttatt cttgtatgta ggtgcttcag aaagcaaaac 180
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<210> 361
<211> 1221
<212> DNA
<213> Homo sapiens

<400> 361
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<210> 362
<211> 684
<212> DNA
<213> Homo sapiens

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<223> n = a,t,c or g

<400> 362

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atgaagcctt acccagggac agaggaagaa agcttcatca cccctctggt gctctattgt    180
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<210> 363
<211> 933
<212> DNA
<213> Homo sapiens

<220>
<221> misc_feature
<222> (1)...(933)
<223> n = a,t,c or g

<400> 363

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<210> 364
<211> 777
<212> DNA
<213> Homo sapiens

<220>
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<222> (1)...(777)
<223> n = a,t,c or g

<400> 364

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actacgatga cttgtctccc acggtcatcc gctcagggct gaaaggtaca gaatgctgca    180
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ctcaaaggga gattcctgac attcagcctt tgtcttagcc tgctcttttc ctcactatga    420
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ctgtgcgata aagttacctt gaaagacttg gggagggtct gcaattacta gactgaggct    660
ttgttgtgaa gggcaccaat caaggggctg ataccttct tgataaaaat tatggagggg     720
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<210> 365
<211> 1157
<212> DNA
<213> Homo sapiens

<400> 365

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agatgtctat agactgtttt atcatatggt agtgttttat catatatggt aggattttac    180
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tatttttatt ggcttccttt agatttaaag aacatgtaca gaataatttg cctagagatc    360
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gtggtgatga tggtcctcct ggtgatgatc taccatcggg aattgaagac ataaccgatc 480
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<210> 366
<211> 1158
<212> DNA
<213> Homo sapiens

<400> 366
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<211> 963
<212> DNA
<213> Homo sapiens

<400> 367
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<211> 842
<212> DNA
<213> Homo sapiens

<400> 368
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<213> Homo sapiens

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<212> DNA
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<211> 538
<212> DNA
<213> Homo sapiens

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attgaacaga agcttccggt aaataagggc cccgtcggca agacagcata ctgctgtcac    180
aagtgcaaac acccctccac caactgtcaa tgttgtggtt tctggtatca gtgccaacac    240
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ctgagtaggt tctgcgaagc gatagcaacc gccaccgcgg cggagcaccg ccctccccta    480
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<211> 1209
<212> DNA
<213> Homo sapiens

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<223> n = a,t,c or g

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ggattacagg cgcctgccac catgcccagc tgatttttgt atttttggta gagacggggt    180
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ttttaatgag ttaatccatg caaagatctt ctaacagtgc caggcacatt gtaaaatgtt    420
aactacgctg ttactattat tacacaaaag gatctttaga ggaaactttc acattctaca    480
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<210> 374
<211> 1083
<212> DNA
<213> Homo sapiens

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aaa 1083

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<211> 710
<212> DNA
<213> Homo sapiens

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gtccttgacg actg 374

<210> 377
<211> 396
<212> DNA
<213> Homo sapiens

<400> 377
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tgacagatct taccaaagtc cacacggaat gctgccatgg agatctgctt gaatgtgctg 180
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<210> 378
<211> 638
<212> DNA
<213> Homo sapiens

<400> 378
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<210> 379
<211> 3043
<212> DNA
<213> Homo sapiens

<400> 379
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<211> 497
<212> DNA
<213> Homo sapiens

<220>
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<222> (1)...(497)
<223> n = a,t,c or g

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gagccaaaga ggccgaccca agtgggagaa ggtctctcgg aagcccaggc ctcgagtgtg 240
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cactggggtg ataccctcgt agtgtgaagg gaacagggca gattcagaga ctgcagcacc 360
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agatgaggtt tatcctg 497

<210> 381
<211> 777
<212> DNA
<213> Homo sapiens

<400> 381
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<210> 382
<211> 659
<212> DNA
<213> Homo sapiens

<400> 382
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<210> 383
<211> 392
<212> DNA
<213> Homo sapiens

<400> 383
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catctctaga attgaattta tttgtttgtg tgtttgtgtt tttttcaggg tgatttggtt 180
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gccaattatc ttgcctctcc acccttagtg gtagcttatg ccatagcagg cacagtgaat 300
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<210> 384
<211> 853
<212> DNA
<213> Homo sapiens

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tatactctct tcc 853

<210> 385
<211> 965

<212> DNA
<213> Homo sapiens

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catctaaaat gaagtaagca ttttagagct aaattagaga agggataatt ccccattttt 180
cattccatgc ctcactctgt ccttctttat gcccaatgtc cctgaatcca gaatttctct 240
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<210> 386
<211> 422
<212> DNA
<213> Homo sapiens

<400> 386
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gc 422

<210> 387
<211> 435
<212> DNA
<213> Homo sapiens

<400> 387
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cctcaaacta cttaactgtg tatctgcctg gaatatgaat ataagactga aatgtctgtt 180
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aaaaattagc tgggg 435

<210> 388
<211> 473
<212> DNA
<213> Homo sapiens

<220>
<221> misc_feature
<222> (1)...(473)
<223> n = a,t,c or g

<400> 388
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gtggccagcg cctcctcctc ctcccagccc tgaggaacca tccttgtcct caaggtggaa 180
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agatgggaga gcccaaaacc tcccgccacc cagtcctcct tnngccccac gaaatcgtcg 420
ncccggggnt tccggngang gngtccaatc gaacggcttc aatggagcca cac 473

<210> 389
<211> 376
<212> DNA
<213> Homo sapiens

<400> 389
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tgggggaggg ccaggcacca tgaagccagt gtgggtcgcc acccttctgt ggatgctact 120
gctggtgccc aggctggggg ccgcccggaa ggggtcccca gaagaggcct ccttctacta 180
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agaagggaat cgaattctac agtgatctta tcgatgccct tctgagcagc aacatcactc 360
ccatcgtgac cttgca 376

<210> 390
<211> 906
<212> DNA
<213> Homo sapiens

<400> 390
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<210> 391
<211> 680
<212> DNA
<213> Homo sapiens

<220>
<221> misc_feature
<222> (1)...(680)
<223> n = a,t,c or g

<400> 391
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gatgcgtggc cttctggcca ggcgtctgcg aaatcatatg gctgtagcat tcgtgctatc 180
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agacactgtn cgggggccca 680

<210> 392
<211> 1983
<212> DNA
<213> Homo sapiens

<400> 392
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agg 1983

<210> 393
<211> 859
<212> DNA
<213> Homo sapiens

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<210> 394
<211> 1407
<212> DNA
<213> Homo sapiens

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tcatgtacag gtttttatac ttcccagcta ttccatctgt ggatgaaagt aacaatgttg    660
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caaatttgtt cagtaaatcc atagatgact acattttgag aaatactaag ataatactaa   1140
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gcaagctctg cttcccgggt tcacaccatt ctcctgcctc agcatcctga gtagctggga   1320
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<210> 395
<211> 319
<212> DNA
<213> Homo sapiens

<220>
<221> misc_feature
<222> (1)...(319)
<223> n = a,t,c or g

<400> 395

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gagaaaacta agcaaagctg tgtggaagag cagaggcagg aagaaaaaaa caaagacaat    180
attggttttc accatttggg caagagaata aatcaagagc tatcatctaa agaaaatatt    240
gtccaggaaa gaaagaaaga tttgtccctt tctgaagcca gtgagaataa gggaagtagt    300
aaatctcaaa attatttcn                                                 319
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<210> 396
<211> 2704
<212> DNA
<213> Homo sapiens

<400> 396

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gtcatgggtg tctcaatgac agtaacgttc ccgaaccccg gaccttagct gtcatttcac    180
ctgcgtcgtc ccggacgcca tttggctgtt gacgtggttc cgagccagca aataacgcca    240
gcagccctcc cagatccacg ccggcccgtc tctccgccgg ccccctcctc gcagtggttt    300
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ccgtctcctc tgggtggccc cagtgcgcgg gctgacactc attcagccgg ggaaggtgag    420
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gccgggcaga caaaagaggc cgcccgcgta ggaaggcacg gccggcggcg gcggagcgca    540
gcgatggccg ggcgaggggg cagcgcgctg ctggctctgt gcggggcact ggctgcctgc    600
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aaaa                                                                2704
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<210> 397
<211> 1743
<212> DNA
<213> Homo sapiens

<220>
<221> misc_feature
<222> (1)...(1743)
<223> n = a,t,c or g

<400> 397

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tggccccaaa ggagctcctt aagccaaagt acttggtaca aagagaccca tattcctata    180
aacatgttaa gtttgttcct aagcattcca gacttttaga ataagaactt catttccaac    240
ttttttattt attaacatgg ggctaaactt ttaagaaaca accctaggtc ttctatttcc    300
caggagctgg ttcaaagtct taaatgacaa tataacttca ttatgaaaat atactgaaaa    360
ggtacaaggg gctgatgtaa aaacggttaa tcaagggttc ccaggcatcc atggggactt    420
aagggtaacc tgaaagaata acccccagcc caggctgcaa ccagccaggc caggatgtgc    480
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aag 1743

<210> 398
<211> 315
<212> DNA
<213> Homo sapiens

<400> 398
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gctcggtgtt agagcggtcc ctctgtgctc tgcctggcag ggcgctgttg gcctggtctc 180
cctcactatt tctatttgca agcatgggct ttctttccag cagaatctgg ttcctgggaa 240
gagtaatgtt ccaaaggcct ctgatatgcc tcgatgccct cctgtcgacg cggccgcgaa 300
ttccagatct atgaa 315

<210> 399
<211> 397
<212> DNA
<213> Homo sapiens

<400> 399
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gagcagcata attcatgttc tgttttccta caaatccagt ctttgacttc aaggttataa 240
gccacagaaa atactctgtg agtgatgacg tggggaatgt gtttggatag gatcactagg 300
gatgcaggca acaaaggaca atgacacatg ctttggggtt tctgtgtttg ttttttttcc 360
agcgatgagc tactcctggg tcatgagaag gcccctg 397

<210> 400

<211> 4175
<212> DNA
<213> Homo sapiens

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<210> 401
<211> 1703
<212> DNA
<213> Homo sapiens

<220>
<221> misc_feature
<222> (1)...(1703)
<223> n = a,t,c or g

<400> 401
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<210> 402
<211> 1433
<212> DNA
<213> Homo sapiens

<220>
<221> misc_feature
<222> (1)...(1433)
<223> n = a,t,c or g

<400> 402
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<212> DNA
<213> Homo sapiens

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<212> DNA
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<213> Homo sapiens

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<212> DNA
<213> Homo sapiens

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<212> DNA
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<212> DNA
<213> Homo sapiens

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<212> DNA
<213> Homo sapiens

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<211> 1063
<212> DNA
<213> Homo sapiens

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<212> DNA
<213> Homo sapiens

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<213> Homo sapiens

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<212> DNA
<213> Homo sapiens

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<211> 1509
<212> DNA
<213> Homo sapiens

<400> 450
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<211> 878
<212> DNA
<213> Homo sapiens

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<211> 4710
<212> DNA
<213> Homo sapiens

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<212> DNA
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<400> 453

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<212> DNA
<213> Homo sapiens

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<212> DNA
<213> Homo sapiens

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<211> 1322
<212> DNA
<213> Homo sapiens

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<211> 1842
<212> DNA
<213> Homo sapiens

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<212> DNA
<213> Homo sapiens

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<210> 460
<211> 620
<212> DNA
<213> Homo sapiens

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<211> 1477
<212> DNA
<213> Homo sapiens

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<211> 458
<212> DNA
<213> Homo sapiens

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<223> n = a,t,c or g

<400> 462
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<210> 463
<211> 1280
<212> DNA
<213> Homo sapiens

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<211> 2290
<212> DNA
<213> Homo sapiens

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<212> DNA
<213> Homo sapiens

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<213> Homo sapiens

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<212> DNA
<213> Homo sapiens

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<223> n = a,t,c or g

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<212> DNA
<213> Homo sapiens

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<211> 1103
<212> DNA
<213> Homo sapiens

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<212> DNA
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<212> DNA
<213> Homo sapiens

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<212> DNA
<213> Homo sapiens

<400> 484

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<211> 478
<212> DNA
<213> Homo sapiens

<220>
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<222> (1)...(478)
<223> n = a,t,c or g

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<210> 486
<211> 477
<212> DNA
<213> Homo sapiens

<400> 486
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<210> 487
<211> 4198
<212> DNA
<213> Homo sapiens

<400> 487
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<210> 488
<211> 861
<212> DNA
<213> Homo sapiens

<400> 488

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<210> 489
<211> 848
<212> DNA
<213> Homo sapiens

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<223> n = a,t,c or g

<400> 489

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aactgagggt aggaagtgga aaacaacctg ggcacatgga aaaaccccat cctactaaaa 660
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cgctttagaa gcaactcttc cccgggcggg ggatttaaat tttttaaggg accaaaataa 840
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<210> 490
<211> 1621
<212> DNA
<213> Homo sapiens

<400> 490
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<210> 491
<211> 466
<212> DNA
<213> Homo sapiens

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<210> 492
<211> 767
<212> DNA
<213> Homo sapiens

<400> 492
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<210> 493
<211> 852
<212> DNA
<213> Homo sapiens

<400> 493
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gaaacacctt gg 852

<210> 494
<211> 849
<212> DNA
<213> Homo sapiens

<220>

<221> misc_feature
<222> (1)...(849)
<223> n = a,t,c or g

<400> 494
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ggctgaggga ggagaatctc ttggaccctt ggaggggga gggtccattt aaccaaaatt 780
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aaaaaaaac 849

<210> 495
<211> 950
<212> DNA
<213> Homo sapiens

<400> 495
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ctggagagtt tattgggagc ccaggaatat tcatttttaa tacacacaca cacacacaca 180
cacacacaca cacactgatc agagtaacag gagtttctct caggagtcat actccatgag 240
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gatacccatg tctctattaa cagctttatc tctttctaca gttttgggta tttgataagg 540
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atttttttct tgaaactcta cagtatacta caatagtgaa ggaaactatt aacatgagag 720
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ttaaaattca ttatgaaaat tagatttatt ttaaatactt tcaagtgtat acatttttat 840
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<210> 496
<211> 838
<212> DNA
<213> Homo sapiens

<400> 496
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ttagcttgcc atggaatatt caagaactat cacatacgtg tggaatacag cgcggatccc 720
gccttaataa ctaactttgg tgggcccggg gggggatcat aagaaaggct ttaaaacctt 780
tggccaacat gagaatcccc tctctagaga atagagagtt acctccgacg cgccgcgc 838

<210> 497
<211> 598
<212> DNA
<213> Homo sapiens

<400> 497
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accaggcaac ttcgtgtgca gcaatggacg gtgcatcccg ggcgcctggc agtgtgacgg 180
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ctgcgatgga cagaataact gtcaagacaa cagtgatgag gaaagctgtg aaagttctca 480
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<210> 498
<211> 1902
<212> DNA
<213> Homo sapiens

<400> 498
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atgttgcgac actcctgctc cttgagcttc acgaaggcat agaagacacc aaagtggaac 180
tggttcagga aggccaactt gttcagcttt acctcgtgct caaagaatcg gtcctccagc 240
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cactttcatg ttcctcccta actccctgac ctgagaaccc tggggcctgg gggcagtttg   1740
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tgtgtgtcct tggggcctgg ggggatgttg ctcctcagct ccctccctca gccctgcccc   1860
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<210> 499
<211> 2122
<212> DNA
<213> Homo sapiens

<220>
<221> misc_feature
<222> (1)...(2122)
<223> n = a,t,c or g

<400> 499

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accatgctca gataattttt taactttttg tagagaaagg gtctcactat gttccccagg    180
ctggtctcaa gcgatcctcc catctcagtc tcccaaagtg ctgggattac aggcatgagc    240
caccactgtg cctggcctaa aaattttttg ttaaaaatgc tttccaccgg ccgggtgcag    300
tggctcatgc ctataatttt tttgtttttt cagaagatgg gaggcaacat ggtaggttca    360
caattaaaat tgtcttgaaa gtatttattg tttaataatt ctttctcccc tcagccccat    420
ccggccactc tctctttctg cttttctgat catcctaaag gctgaataca tcctcctcat    480
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cacgtcactg tggggtagag tgctagtttt caacaaatgg tgggtgttct tatgggctcc    600
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aagaatcctg ccaaatctta cggtgttttg aacagtcaaa tatttctctt gtaattcagg   1620
ctcactgccc aaaggcaaga gaacttcaat ataactgtcc ttcattctcc taggaggcag   1680
tccatcctgt gatttagcca agaaactatg aagtaatttc ctttcttcca ttgccttcac   1740
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atggtctctc cagtttgtgg atgctcctac tatctcccgc aacttatctc gaacttcatg   1800
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ctgccgtgcg cgcgatggag aaccgggccc cgcgcgctag tcggcggagg gaaactgagg   1980
cgataaaaga cgcacgagta ccagaccgcg cccttgctga ggacagcccg ggagccggac   2040
agcggcccgg ctcgagcggc cgctcgagcc gggaattcca ccgcnctcct ataatggtct   2100
tctatggggg gggggggggg cg                                            2122
```

<210> 500
<211> 458
<212> DNA
<213> Homo sapiens

<220>
<221> misc_feature
<222> (1)...(458)
<223> n = a,t,c or g

<400> 500

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tgacctgcgt cggccactgc agcaatggcg gctcctgtac catgaacagc aaaatgatgc    180
ctgagtgcca gtgcccaccc cacatgacag ggtcccggtg tgaggagcac gtcttcagcc    240
agcagcagcc aggacatata gcctccatcc taatccctct gctgttgctg ctgctgctgg    300
ttctggcggc cggagtggta ttctggtata agcggcgagt acaaggtgct aaaggcttcc    360
atcaccaacg gatgaccaac ggggccatga acgtggagat tggaaacccc acctacaaga    420
tgtacgaagg cggagagcct gatgatgtgg gaggccta                            458
```

<210> 501
<211> 511
<212> DNA
<213> Homo sapiens

<400> 501

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gcctttcttt tatacatctt cctcaaccta cagctcatga tcttgcaggt ccttcacctt     60
tactggggtt attacatctt gaagatgctc aacagatgta tattcatgaa gagcatccag    120
gatgtgagga gtgatgacga ggattatgaa gaggaagagg aagaggaaga agaagaggct    180
accaaaggca aagagatgga ttgtttaaag aacggcctcg ggctgagag gcacctcatt     240
cccaatggcc agcatggcca ttagctggaa gcctacagga ctcccatggc acagcatgct    300
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ctctttctca ccaccgcctt ccctcccacc taagatgtgt ttaccaaaat gttgttaact    420
tgtgttaaaa tgttaaatat aagcatgccc atggattttt actgcagtta ggactcagac    480
tggtcaaaga tttcaaagat ttctccacaa a                                   511
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<210> 502
<211> 964
<212> DNA
<213> Homo sapiens

<400> 502

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ggaaagcccg cctcctccct cggccggccc tggggccgtg tccgccgggc aactccagcc   180
gaggcctggg cttctgcctg caggtgtctg cggcgaggcc cctagggtac agcccgattt   240
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cacgtcctgc ttcaggagga ggtggccgag ctgcagggcc aggtccagcg caccgaagtg   420
gcccgcgggc ggctggaaaa gcgcaattct gacctctttg ctgttgttgg acacgcacaa   480
gaaacagatc gaccagaagg aggccgacta cggccgcctc agcagccggc tgcaggccag   540
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agaatatgaa agttttcagt gtggacagca gatgaaggaa ttgagagcac agcatgaaga   780
aaatattaaa aagttagcag accagttttt agaggaacaa aagcaagaga cccaaaagat   840
tcaatcaaat gatggaaagg aattggatat aaacaatcaa gtagtaccta aaaatattcc   900
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tggg                                                                964
```

<210> 503
<211> 681
<212> DNA
<213> Homo sapiens

<400> 503

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tgcggcgccg cgacaggaag cggcgggcga gccgagtgtc cttgcgcgtg gatccgagcg   180
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ctccggactg ggaccccact ccgagggcag ctcccggcct tgccggccca ataaaggact   660
tcagaagtga aaaaaaaaaa a                                             681
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<210> 504
<211> 4179
<212> DNA
<213> Homo sapiens

<400> 504

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<211> 971
<212> DNA
<213> Homo sapiens

<400> 512
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<211> 422
<212> DNA
<213> Homo sapiens
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<211> 1568
<212> DNA
<213> Homo sapiens
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<210> 515
<211> 857
<212> DNA
<213> Homo sapiens

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<210> 516
<211> 2133
<212> DNA
<213> Homo sapiens

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taagttcact aaaaccctgg attgaaagtg aaaattatgt tcctagaaca aaatgtacaa   1140
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<211> 1404
<212> DNA
<213> Homo sapiens

<400> 517

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<211> 698

<212> DNA
<213> Homo sapiens

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<210> 519
<211> 752
<212> DNA
<213> Homo sapiens

<400> 519
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<210> 520
<211> 2533
<212> DNA
<213> Homo sapiens

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<210> 521
<211> 545
<212> DNA
<213> Homo sapiens

<400> 521
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<213> Homo sapiens

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<212> DNA
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<211> 431
<212> DNA
<213> Homo sapiens

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<210> 532
<211> 2053
<212> DNA
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<212> DNA
<213> Homo sapiens

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aaaaaaa 1567

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<211> 345
<212> DNA
<213> Homo sapiens

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<211> 781
<212> DNA
<213> Homo sapiens

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<211> 590
<212> DNA
<213> Homo sapiens

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<211> 442
<212> DNA
<213> Homo sapiens

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<211> 901
<212> DNA
<213> Homo sapiens

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<211> 384
<212> DNA
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<211> 732
<212> DNA
<213> Homo sapiens

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<223> n = a,t,c or g

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<211> 842
<212> DNA
<213> Homo sapiens

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gg 842

<210> 543
<211> 1100
<212> DNA
<213> Homo sapiens

<400> 543
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<210> 544
<211> 939
<212> DNA
<213> Homo sapiens

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tttgggagaa cgcggtctac tttgtctgca attattctcc aaaggggaac tggattggag 840
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<210> 545
<211> 1053
<212> DNA
<213> Homo sapiens

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gaatctctct cgaaaagctt cctttggaaa ttt 1053

<210> 546
<211> 715
<212> DNA
<213> Homo sapiens

<400> 546
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ggtacgcaga atacattatc cagcttaaac tcaacaaacc ctgtttcaac aaactgaaga 420

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<210> 547
<211> 812
<212> DNA
<213> Homo sapiens

<220>
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<222> (1)...(812)
<223> n = a,t,c or g


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ccgagcaacg gcagcacata ttttctttca aagtacaaat atatcattac aaaaactgac     180
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<210> 548
<211> 578
<212> DNA
<213> Homo sapiens


<400> 548
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tacttcattg tccagtgttt ctcccatttt actacactgg caaatgtgtt tatggaggaa     420
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<210> 549
<211> 428
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<212> DNA
<213> Homo sapiens

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tttggggtgg acagatgccc actgaagagc tttggaagtc aaagaagcat tcagtgatgt 360
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ctctttgc 428

<210> 550
<211> 849
<212> DNA
<213> Homo sapiens

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gctcggagaa cgacgccctg tatgagtacc tgcgccagag cgtgggcaac gaggccgaga 480
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aaaaaaaaa 849

<210> 551
<211> 648
<212> DNA
<213> Homo sapiens

<220>
<221> misc_feature
<222> (1)...(648)
<223> n = a,t,c or g

<400> 551
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aacggcccag gggacgccct cggcctcgag gcgggggggg ccccggaccg ccccccccacg 420
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cgaatgcccg gcggccgcat gacccccgcc ccagaggctg ctcgttcttt tgaacaaggc 600
acgcgcccta ttaattctcc ctgtccgggg gaccggtccg atcgaacc 648

<210> 552
<211> 713
<212> DNA
<213> Homo sapiens

<400> 552
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gcttgtttta agtacctgtc actgactcat tccccaactg aagcctaacc ttccttttt 660
gtggggaggc acacctcagg ggtagctgcc attcatcctt tcttcctgag gcg 713

<210> 553
<211> 714
<212> DNA
<213> Homo sapiens

<400> 553
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aatttctact ctttccgagg ggctaatgct gttctcagct gtgaaacttt attgttgtca 180
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<210> 554
<211> 836
<212> DNA
<213> Homo sapiens

<400> 554

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aataaatgtt tagcgcgaga tagtccagtg taaccatgaa ttcaaaattg ggtgaaatga 180
gaaggcaaat agcatgtcag gcagtcaggt tatctcagag tgggggacat tgatggagag 240
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<210> 555
<211> 1765
<212> DNA
<213> Homo sapiens

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gttcagagcc cttgagctag catttcatta tgaccgtgat tttttccccg caccactttc 420
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<210> 556
<211> 1044

<212> DNA
<213> Homo sapiens

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<210> 557
<211> 1372
<212> DNA
<213> Homo sapiens

<400> 557
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agtatacttt ctctacaaag ctctactttt tgaggcttca aataaattac atttatccta    720
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gcctgtaatc ccagcacttt gagagagcaa ggtgggtgga tcacttgagg tcaggagttc    960
aagaacaggc ctggccaaaa gggggaaacc tggtctttaa taaaaaggcc aaaattaacc   1020
agggcttggg ggcaggcccc ctgtgttccc agctggcttg ggaaggcctg gcgcccagga   1080
aaaaatgct                                                           1089
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<210> 582
<211> 443
<212> DNA
<213> Homo sapiens

<400> 582

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cgggtcgacc cacgcgtccg gagcgccccg gggagctcgg agcgcgtgca cgcttggcag     60
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aaaaaagtgg aaaatgggag gcatgaaata catcttttcg ttgttgttct ttcttttgct    180
agaaggaggc aaaacagagc aagtaaaaca ttcagagaca tattgcatgt ttcaagacaa    240
gaagtacaga gtgggtgaga gatggcatcc ttacctggaa ccttatgggt tggtttactg    300
cgtgaactgc atctgctcag agaatgggaa tgtgctttgc agccgagtca gatgtccaaa    360
tgttcattgc ctttctcctg tgcatattcc tcatctgtgc tgccctcgct gcccagaaga    420
ctccttaccc tcagtgaaca atg                                            443
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<210> 583
<211> 2590
<212> DNA
<213> Homo sapiens

<400> 583

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tttttttttt ttgtataaaa acggcatatt ctttattttg catactttaa tttcagaaca     60
aaatgaagaa aataaaataa accacaatac acaacatcca atcctgctgt caagagtaga    120
gagggaatgg ggcttgacac ccttagttta ctgccttcaa cacaaggaca ggagagggaa    180
aaaaacacta gacaccagca gggggagcca ggtgggacag gggcactcga ggctgcagtg    240
ggagcccatg ggacactat acaagggcac aagttttcca actatgaact cctaacctaa    300
tcgacttctt ccatgcgaga cgcatcctca tcgccctcga gagggggat ctcatcagga    360
actgcagcat tgggttcctc tgctgccact tcatcttcat caatacctag acctagcttg    420
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agcagggcgg tttcaaacag cagcaccacc aggtccttaa ctgccttatc attcttgtcg    540
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agtctattgg agattgtcac cttctcaacc ttcttatcta agatttcttt catgagcttg    780
cagaggttct caaactttgc cttgctctct tccatcttct tcttctcctc ctcatcctca    840
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cgcactcgct ccacaaaagc tgagttggcc acctgctctt tgctctcacc agtgatgtaa   1020
tagatggact tctgtgtctc cttcatgcga gaaacatact ctgacagaga tgtcatctca   1080
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tccttgtctt ctgccagctc agagaagagc tcaaggcact tcttaacaat gtttttgcga   1260
atgactttca agattttgct ctgctggagc atttctcggg agatgttcag gggcagatcc   1320
tcagagtcaa ccacaccacg gataaaattg agatactctg gtatcaactc atcacagctg   1380
tccatgatga acacacggcg gacatagagt ttgatgttgt tctttttctt cttgttctca   1440
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gaacctccag cagaagactc ccaagcatac tgttcatcat cgttgtgctt tgtgatcaca   2040
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gttcccaaat tatttatgag atcagctttg gtcatgccaa tgcctgtgtc taccaaagtc   2220
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ttcgaagggt ctgtcaggct ctcatagcga atcttgtcca aggcatcaga agcattagag   2340
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agttgggcaa tttctgcctg aaaggcaaaa gtctccacct cctcctctcc atggtgcact   2460
tcctcaggca tcttgaaaag aaaaggatta tacgtaatag tgagcaacgt aggcttgctt   2520
tccgataccc agacagtccc aacactgcgc cggagtgact agagagagat actgcgtgcc   2580
ccaagtcgcc                                                          2590
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<210> 584
<211> 425
<212> DNA
<213> Homo sapiens

<400> 584
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tgccttagga cagtttttc taattatggg taacacgcag aggtgtgatg actttcctac    120
tgaaagtccc ccagcaaaga caaacgtttc ccgcgcaggc ttgtcccctc cgtgtgaggc    180
cctacatggt gtagaaagta ggggcagctg cagccacggg aagctgcaaa gccctcctgg    240
gagagactgg ccgcagggtg acccacagga caggcccaag cgcagatggc agaggccagg    300
acctgctggt cggggcgccc cagaccccac tcctaagggc caggggggcag cagtcccacc    360
gcgctctgcc agcatgtttc tgatccacaa gcagatgtgg gcctatggct ttggggactg    420
aaaga                                                                425
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<210> 585
<211> 841
<212> DNA
<213> Homo sapiens

<400> 585
gcagtgcgcg tggaattcat ttcttcccct tatggccaat ttccaggcct ccaggcccct     60
ctctggacct ggaatgaccc tagcatcttg gctcttgctt aaagccattc cagatttcaa    120
gaaataccat ttaaggcaat aagggaccta tttatttctc taatgaggca actggacttc    180
agaaaatgta agtgacttga caagttgcat tcccttagtc attcagctgc cttcctggaa    240
cacataagca aacaatcctc aatgtaatgt cagagattgg taagtgcttt gagaaaacac    300
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tagagcaagg taagaaaatg acagagcagg gagtctattt taaataaggc agtagagaaa 360
gccctggtac agcaggtggt agtcacatga agttatgggg agggggttcc aggaagaggg 420
aagagcaaat aacaaggacc tggaggtggg aattagctga atgaacaaaa cacaaagcaa 480
taagaaatgg aattagagag gaagacagag cccagatcat ttaagctttg aaggccaagc 540
tccgactttg gactttattt gaaagtgtct gtaaagcttt taaagagtct taaaactctt 600
ggccaggcgc gggggctcat gcctgtaatc ccagcacttt gtgaggccaa ggcgggctga 660
acacaaagtc aggagttcga gaccagccgg accacatggg ggaaccccat ctttactaaa 720
aacacaaaca ttagctgggc atggggggat gcacctgtaa tccccactac ttaggaggct 780
gaggcaagag aatcgctttg acttccagag gggggagttg ccattcgccg aaaaacaacc 840
c 841

<210> 586
<211> 787
<212> DNA
<213> Homo sapiens

<400> 586
aagggtctag aaagaatggg ctccccctgg gtgctgcatg cctctggggt gagcacagtc 60
ctggccctca tgagcccacg cagagagcgt ggcaatcctg tgtctcctgc aggtatgcag 120
gcggcccggg gggcctgggc ctcccctcac atgctgcaag accctccact gacttcacgc 180
aagcggcagc tgctgcagct gtggctgctg cggcagccac tgccaccgcc acagccacag 240
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gccccctcag cagctcctcc cacatggcag ccagcctgag ggcctgggag gaggtactca 360
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agagagatgt gtgggaagga gtgagggtct gaggggagag gtttctgggg tgtcctgtga 480
aagggtatgg tgcccacatg ggtgggtggg cgggttttat gcctatcttt tggagccctt 540
tgtgggtggg acctggacca ttcttctttt ttctcttcct agatgggggc cggacagtct 600
tttaacagcc agtttctgca gcatggaggt ccccgggggc ctagtgtccc cgctggcatg 660
aaccctactg gcataggagg ggtaatgggc ccctctggcc tctccccctt ggctatgaac 720
cccacccggg cagcaggaat gacacccttg tatgcagggc agcgtttgcc ccaccatggg 780
tatcctg 787

<210> 587
<211> 363
<212> DNA
<213> Homo sapiens

<400> 587
ctgactcact tacatggcat ggactatacc cgtgactaca cgagatgcat ggtcctatca 60
tggctgacct tgatcgaagc tctcgctgat gtcatgacta ccgatggcaa catgcttcaa 120
ctgttctgtg ttgagcgtac taacctactc gtcaatcaga tacggatgac cttgtatgct 180
caataccgac acgtccgacc cttccgcaca atcatgaagc ccatcttgac ccgagaggtg 240
cagacaaagg actagtcgga cccggccaat ctggtgactc cccaccgcct tggactacac 300
gtcttaaagg cttgccaatc tatttatcct ctccatgatg tcttcgttag aaaagtagac 360
atg 363

<210> 588
<211> 814
<212> DNA
<213> Homo sapiens

<400> 588
gtggaattcc ccccacaggc tccttgtcat gcgaggttgc agtctgattt tcatctactc 60
agattaaatt taatcttgaa gatatagtag aggactggaa tgaggatctg tgactatggg 120
tggctttatt ttcttctttt gacacttgtt tattttctgt aatgagcatg ggtagcttat 180
gattaacaaa cattaaattg gatattcttg aaaacagcaa aaacattttt aatgaaatgg 240
catgctaatc tcattaattt cattattttg tgataaagtc taatgatgag atgagagttg 300
taaactaaga gacgagtggt aatccttggc accctttctt attatgctat ttatttgact 360
tggagagttt tacttgtctg tttttagaga gtatgttaat tgagtgctca gtatgcatta 420
cgaataatct tgtctgtttt cttgtggaga ttctgaaggc ccttttgctc tttctgtaaa 480
agccaagcag actgtattaa cttctgggtt aatttgaaaa atgaatgtgg aacttgttgg 540
cacaacacct taaagaattg catgtttaat aactggaagg ctttccatta gatttggctc 600
tagcctgaat taataatgat gctgacttat tgggaataga agaccccgcc cttggaccgc 660
ctaggaccaa agaaatgggg cctggtctgc aaacccgtcc tgcccccctt gacccgggcc 720
cccctccgct ctgggaacga cactcaccgc ccccgcgacc gaacttgtca tctacaaacc 780
ccgcgcgccc tccgcccacc tcacccacag gacg 814

<210> 589
<211> 794
<212> DNA
<213> Homo sapiens

<400> 589
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aaaaccatta cttcccactt ggagagattt ttgaggatta aatgagataa tgcatgaaag 120
ccctctagct tgggcatcag tacacctgag ttcccttccc ttgctctgca cagcctgctc 180
atcaccactg atggggaact ctgtcctctg tagggcccct gcagacatgg gccttgcctg 240
gatgctgctg ctgtcggagc ctaggagagt tgtgcctggc atcgcagcac aggtactcac 300
agctctcaga aggagactcc tgtctgggac cctgccctca ttcccacgta ggaaaaatcc 360
tttacatgag catctcctgg ccttcattgt taggttgtag actacaatga atgatattct 420
gtgtttaatt acattatgca caacactcta cagagtgggt ggttttgaat cccaaccact 480
aatttacgaa gtggagcggc tctgctggct ctgtgaagta tgtgttgtgg agccagaggt 540
gatgctgttg gatgtgggtg gtgatttacg ggagagcagc ataagcagag gaaggcacag 600
agacctgggt tcaaatccca ctgccagggc tatctgacgt gagacttcgg acaagttatt 660
taaccttaaa gcttagtggc cttgcatgta aaaaacaaat aatgccgacc tcattggatc 720
cttgtggagg agcccctggg ataatggggg gtaccatgca tcagggatca tttccctttc 780
ccttgataaa tgag 794

<210> 590
<211> 1012
<212> DNA
<213> Homo sapiens

<400> 590
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gtttacaaat gccacggacc caaggttccg atccgcgcca aggcgttccg gtcagcagca 120
gccccgcggc tcctcgggcc ccgccgcgct ggcaagcccc agtccccgcc agcccaatcg 180
tgctggcgct ttaaggacgg gcggggcggg ctgggcgaca gcgctggaca cctggagctg 240
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gcatgacagt cagcggcaca gtggttctgg tggccgggac gctctgcttc gcttggtgga 360
gcgaagggga tgcaaccgcc cagcctggcc agctggcccc acccacggag tatccggtgc 420
ctgagggccc cagccccctg ctcaggtccg tcagcttcgt ctgctgcggt gcaggtggcc 480
tgctgctgct cattggcctg ctgtggtccg tcaaggccag catcccaggg ccacctcgat 540

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gggaccccta tcacctctcc agagacctgt actacctcac tgtggagtcc tcagagaagg    600
agagctgcag gacccccaaa gtggttgaca tccccgacta acgaggaagc cgtgagcttc    660
ccagtggccg aggggccccc aacaccacct gcatacccta cggaggaagc cctggagcca    720
agtggatcga gggatgccct gctcagcacc cagcccgcct ggcctccacc cagctatgag    780
agcatcagcc ttgctcttga tgccgtttct gcagagacga caccgagtgc cacacgctcc    840
tgctcaggcc tggttcagac tgcacgggga agaaagtaaa ggcttcctag caggtcctga    900
aaccaaaaga caaaaaaggc tgtgcccttc tcccaaaacc ttaggccggg cgctgggaca    960
acaggaggcc cttcctgcaa acgttcgttg gtgaaaggct ggtcatattt aa           1012


<210> 591
<211> 860
<212> DNA
<213> Homo sapiens


<400> 591
ctccgtgtgg tggaattctt cacatttcag gaagggagac ttggggcctg gagaagcgat     60
gtgatttttc ttttctagtt cagcgctggt tttgatggct ttttatcatg accttgttat    120
gtcttatttt agtttcggcc catttagtgg atacgacaac agtggcccag ggaggtatgg    180
cagagctgag gcttaaccca gggcctgcgc cctccacggc ctgcactgcc ccacctccag    240
ctccttgccc tgttcctccc tctgcaccgg atcagccccc ggactctggg tcacctccac    300
accagttgac agggcccccc agtccccacc gccaaccacc tggccggcta cttgtcagac    360
agacatgggg gcgtgggcat gggtccccac ccctagcctt tgcctctgtc actctacctg    420
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gtcgcccagg ctggagcgca gtggcgcgat cttggctcgc tgcaacctct gtctccgggg    540
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tgcctggcta atttttgcat ttttggtgga gacagggttt caccatgttg gccaggctgg    660
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ggcgtgagcc acctcaccca gcctggagtg tctcatcttc caccactaaa tgaaacgatg    780
gaccctgaac agaaaaagga acagtggtgg aagaactagc aaagcccaca gccttgagtt    840
tggccgtaag tatcaaggtt                                                 860


<210> 592
<211> 825
<212> DNA
<213> Homo sapiens


<400> 592
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gggacggtag ctcctctctg aagctctcag cagggatggt agctcttctc tgccggcaga    120
tcatctctgc agccttcagt ggagagggta ctcctctctg cagctggtcg tctggtccca    180
tcctgtcatc tgtctgcctt ctttgtcctc tggccgtcct ctgccctgct aagcctgagc    240
ccagggcttt tacggacctc agaggggagg aagtgtgtgc cgactggttc atgggcggcc    300
atgggagggt cgaaagaggc accatgagtc cccactctgg tctgtaggac tggcagcctg    360
gcccccagtc ttcaggccct ccctggcctg aaggtggggc cttactgggg acccaccccc    420
ttctgcccag gaattaatct gccttctgct gccattcacg gccctatgac ttggaccaaa    480
ccccactctg acagaggtca ggcagtggga gcaaacaccc ctgaacctgc atggactagg    540
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gcagaacagc accacttgcc ccagcaactc ctaccctagc ccacatccac gagccaaggc    660
acttccccag gaatccacaa gctgccaggt caccacggga gacgaaggca ccaggacata    720
aaaactgcgg gaccagtaca gcattgtgca tttcaggtct ccaaggttct gacccccccc    780
cccccggatg acctgggaac ttgtaggaat cccccgaggg gaggc                    825
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<210> 593
<211> 867
<212> DNA
<213> Homo sapiens

<400> 593
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tttttaaat ttaataccaa tgtttattag ggcagaaaag aagaggaaaa aaatagagga     60
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ttccttaggg caggcagcct gcccgaactc ctggtctggg agttccagct ccatcaaccc   180
caggtaagat tctggttgtt cccactcttg caaactgatg ggaagacctt tgggaggtgt   240
ctatgcttta agctattggt tttagtgatc tatgcaggtt agtaaaatga agcagtatat   300
atatttgcca tttccaaggc aatctttgat atgcccacag ttcacgaggt ctgaagacat   360
ccatttctgc aatttaaaaa caagtgaaag aagcagcctt gtcttgcttc gacattatcc   420
agcttgttgt ctattaaaat gcttgcgagg ctggtcctga tccccttaca caggatgaat   480
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gaatccgggt ttgcttggtc tatcctgtcg ccgagatctt aggagctgtt tggcctgctt   660
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cacaggcaat gtcttcatcc tgccttggtt cctctgcctc ttgctgagtg aatcctccca   780
gactgagtca gccaacttga aggaagccat gccaggccct gcgcttgttt atgctttgac   840
taacgggact tacggtatga tgctcaa                                       867
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<210> 594
<211> 654
<212> DNA
<213> Homo sapiens

<400> 594
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ctgtgagtgt ggcggaattc agatttttca cttttcttct gagctctggt gctttcagag    60
tggtattttt atattcgaat agttgctagt tgtactttta aaagcgattg atgctggagg   120
tcttctattc caccatctcg ctgatgtcag tcctcaaata ataattttat attttagcaa   180
attattttgg ttttaggatt ttgtgtctac gtgacacaga catgaaaaga gatgtactca   240
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tatttaagcc acatgtttta tacatagact gtcctttaaa gagactagat agttctgtgt   360
gtcagcatat agggacagaa tataactaca cattaataat ttctcaagta tttattttag   420
aagtgtaagt aacctttatt ttaattttg ttatattatg cctctgtaat gcagataaat   480
ttttatcttc aggaaatgga aaattttgtc cagagttcag gggaagatgg tattgtggtg   540
ttttctctgg ggtcactgtt tcaaaatgtt acagaagaaa aggctaatat cattgcttca   600
ggcccttggc cagattccca cagaaggtca ggtaaaccct ccattcctgg taaa         654
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<210> 595
<211> 611
<212> DNA
<213> Homo sapiens

<220>
<221> misc_feature
<222> (1)...(611)
<223> n = a,t,c or g

<400> 595

gcggttttcc tcaccagagt ttgataaatc aggggcaagg aggaagttaa acgggcagat 60
gactgcagag ggtccttcca gttctaacat caacggaagc taactacatt ccccactcaa 120
atcatctctg cacatacagc ccgcaggaag ccctttgaaa tgtatttaac cacctttctc 180
gctctcagaa tgatctcaac aagaacagct ttgctttcct tggagctctg catcaatcta 240
ggaaggctgc tttgtctctt cactacttga gcaggatgga gagatatgag cgggaaagac 300
agataagaaa tctgagaaag ccccacaagg tgggttgata gtgtgaagaa catgggctga 360
agcatccaaa tcttggttca gctacttaca gggtaacctt gagaaagtta cttaaacttg 420
tcagctcgga cgggcgtggt ggctcacgcc tgtaatccca gcacattggg aggccgaggt 480
ggacggatca cgaggtcaga tcgagaccac cctggctaac acggtgaaac cctgtctcta 540
ctaaaaatac aaaaaaatta gctgggcgcc tgtagtccca gctactaagg aggctgagng 600
cggagaatgc c 611

<210> 596
<211> 644
<212> DNA
<213> Homo sapiens

<220>
<221> misc_feature
<222> (1)...(644)
<223> n = a,t,c or g

<400> 596
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<213> Homo sapiens

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<400> 606

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<211> 592
<212> DNA
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<212> DNA
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<213> Homo sapiens

<400> 615

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<210> 616
<211> 1927
<212> DNA
<213> Homo sapiens

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<210> 617
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<212> DNA
<213> Homo sapiens

<400> 617
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tgtggctatt ttaaaaggtg tccagtgtga ggtgcagctg gtggagtctg ggggaggctt 180
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<210> 618
<211> 946
<212> DNA
<213> Homo sapiens

<220>
<221> misc_feature
<222> (1)...(946)
<223> n = a,t,c or g

<400> 618
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<211> 354
<212> DNA
<213> Homo sapiens

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<210> 620
<211> 384
<212> DNA
<213> Homo sapiens

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<210> 621
<211> 873
<212> DNA
<213> Homo sapiens

<400> 621
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<210> 622
<211> 875
<212> DNA
<213> Homo sapiens

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<210> 623
<211> 923
<212> DNA
<213> Homo sapiens

<220>
<221> misc_feature
<222> (1)...(923)
<223> n = a,t,c or g

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<210> 624
<211> 1101
<212> DNA
<213> Homo sapiens

<400> 624
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aaggagcgta gaggcacagc g 1101

<210> 625
<211> 1077
<212> DNA
<213> Homo sapiens

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<210> 626
<211> 1085
<212> DNA
<213> Homo sapiens

<220>
<221> misc_feature
<222> (1)...(1085)
<223> n = a,t,c or g

<400> 626
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ggacg 1085

<210> 627
<211> 838
<212> DNA
<213> Homo sapiens

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<210> 628
<211> 845
<212> DNA
<213> Homo sapiens

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<210> 629
<211> 913
<212> DNA
<213> Homo sapiens

<400> 629
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<210> 630
<211> 812
<212> DNA
<213> Homo sapiens

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tttgggggct ttttttacac gaaggaatat tttggattta agagaggaga ctattggacc 720
ccacgtgaag agacactttt agtgtggggg tgtagtacgg gaacacggag tattatatca 780
tcgcctctac cacgaggaca cctacctcgc gg 812

<210> 631
<211> 760
<212> DNA
<213> Homo sapiens

<400> 631
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ctacaaggga cattttagtg taaggcaaaa atataaaatt atcactcata atgtttttc 180
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aactgtccaa ttttatgtgc attgaaagca aaagctagct gagaaaggaa agcttttctc 480
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<210> 632
<211> 1716
<212> DNA
<213> Homo sapiens

<400> 632
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ggtggacctt ggattttggc gggtgaagag gctaggttgt ttaaggaggt ggggcgcgtt 180
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gttgggccgc gagctgatgc tgcagctgtc ggagtttctg tgcgaggagt tccggaacag 540
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caggaacaat gcaaatggag tggacctgaa ccgcaacttc cctgatctca atacctatat 720
ctactataac gagaagtacg gaggccccaa ccaccacctg ccccttccag acaactggaa 780
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<210> 633
<211> 924
<212> DNA
<213> Homo sapiens

<400> 633
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gttaacggaa ctcctgagag gtggatcctt tgagtttaag gacatgcgtg ttccttcagc 180
gcttgttact ttacatatgc tcctgtgctc tatccccctc tcaggtcgtt tggatagtga 240
tgaacagaaa attcagaatg atatcattga tattttactg acttttacac aaggagttaa 300
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acaaatgttc ttttcataaa aaaa 924

<210> 634
<211> 455
<212> DNA
<213> Homo sapiens

<400> 634
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ttttgaccct ctccctctgc ccttaaaatt gttatttcat ttattcattc tacaaatatt 120
tcctcagcat atgctcaggc actgtgctgt ccactggcac aacaatgtga acttggggga 180
gacaaattat aataaattat taaaagagct ataatggata taaagtgtgt gttctgacag 240

aaaatgggga gaaggtggct atttttgata gcgtgtttaa gatcagcctc tatactggcc 300
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ccacaccttg cagtcccagc tattcgggag gctgaggcgg ggagatggct taagcccagg 420
aggcggaggt tgcagcgacc caagatcgca cgaaa 455

<210> 635
<211> 384
<212> DNA
<213> Homo sapiens

<220>
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<222> (1)...(384)
<223> n = a,t,c or g

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attaaaccga gcatgagggt cctagacaag tgggtcagtg cctttgtgca caaaggtttc 240
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cttaacaaat gcggctgggc cgtattctgc gcagcaccgt gatggttacc gttggtggca 360
gacggatggc cttgggacga tttn 384

<210> 636
<211> 1201
<212> DNA
<213> Homo sapiens

<400> 636
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gtgacactga tggtgctgag ctccccactg gctttggctg gggacaccca accacgtttc 180
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a 1201

<210> 637
<211> 981
<212> DNA
<213> Homo sapiens

<400> 637
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ggaaaccggg tataggagga agagagagat ggaaccatgt aaccaccacg accaagaggc 420
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cttcatttgg acccgcagct g 981

<210> 638
<211> 1421
<212> DNA
<213> Homo sapiens

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<210> 639
<211> 755
<212> DNA
<213> Homo sapiens

<400> 639
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agggctgacg tgcgggaggc aggtttgcaa gtgtgactgc ccacctggct tcaaagccag 180
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ggatgacagg gcaagactcc gtctgaacaa agaaa 755

<210> 640
<211> 1776
<212> DNA
<213> Homo sapiens

<400> 640
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tccaataaat gggatttgct ctctgcaaaa aaaaaa 1776

<210> 641
<211> 418
<212> DNA
<213> Homo sapiens

<220>
<221> misc_feature
<222> (1)...(418)
<223> n = a,t,c or g

<400> 641
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acctccaatc acagcccagc aaggtaactg gacagttaac aaaacagaag ctgacaacat 180
agaaggaccc atagccttga agttctcaca cctttgcctg gaagatcata acagttactg 240
catcaacggt gcttgtgcat tccaccatga gctagagaaa gccatctgca ggtgttttac 300
tggttatact ggagaaaggt gtctaaaatt gaaatcgcct tacaatgtct gttctggaga 360
aagacgacca ctgtgaggcc tttgtgaaga attttcatca aggcatctgt agagatcn 418

<210> 642
<211> 731
<212> DNA
<213> Homo sapiens

<400> 642
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atagtttctg gaaagcctga aatttttaaa aattatactc tcacgtaggg gcatcttatg 180
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tcttgttttt gtggacagag agagatctcc ataatttga gatactctat gttaaactat 360
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<211> 956
<212> DNA
<213> Homo sapiens

<400> 643

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<211> 904
<212> DNA
<213> Homo sapiens

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<211> 943
<212> DNA
<213> Homo sapiens

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<211> 782
<212> DNA
<213> Homo sapiens

<400> 647
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<212> DNA
<213> Homo sapiens

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<210> 649
<211> 886
<212> DNA
<213> Homo sapiens

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<212> DNA
<213> Homo sapiens

<220>
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<223> n = a,t,c or g

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<212> DNA
<213> Homo sapiens

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<211> 743
<212> DNA
<213> Homo sapiens

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<210> 653
<211> 1524
<212> DNA
<213> Homo sapiens

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<211> 711
<212> DNA
<213> Homo sapiens

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<210> 655
<211> 1524
<212> DNA
<213> Homo sapiens

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acgcggtggt gcgggggctc ttcctgctca agctgggact tagcctgctc atgttgttgg 1440
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<210> 656
<211> 993
<212> DNA
<213> Homo sapiens

<400> 656
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cgtgatcgtc ctccaatacg tgtgccccgg cacagaatgc cagctcctcc gcctgcaggc 180
gttcagctcc ccggtgccgg acccgtaccg ctcggaggat gagagctccg ccaggttcgt 240
gccccgctac aatttcaccc gcggcgacct cctgcgcaag gtagacttcg acatcaaggg 300
cgatgacctg atcgtgttcc tgcacatcca gaagaccggg ggcaccactt tcggccgcca 360
cttggtgcgt aacatccagc tggagcagcc gtgcgagtgc cgcgtgggtc agaagaaatg 420
cacttgccac cggccgggta agcgggaaac ctggctcttc tccaggttct ccacgggctg 480
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cggcaagcgc gacgccaggc tgagaccgtc caggtggagg atttttcaca ttctatatgc 600
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gtaccctggt gaagaactgg tctggcttgc ccccttcaa aagattatag gcctggccct 900
tttaatctac ccctaaacca ccccggttgt gccttgtctt tagctacctt ttatatttat 960
ggggtgggtc acactctctt ccaccatctt ccc 993

<210> 657
<211> 969
<212> DNA
<213> Homo sapiens

<400> 657
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taaatttacc attttgctat ttgttttcta tatttctcct gtcttttttg atgttgttat 180
tttctgcatc cttaactggc ttcctttgtg ttaaataaat attttccaat gtagattttt 240
agtttttctc tttttcagct gtatgacatt agtactcttc ctagtgcttg ctctaatgat 300
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aaaatacaga aacttccctt caatattgct tcatttcttt catctttggt tatcattttg 420
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gtaaatcttt ggaggaaatt aatttcaact gaagaaattt taaaaactat ttttgggaag 660
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aaaaaaaaa 969

<210> 658
<211> 572
<212> DNA
<213> Homo sapiens

<220>
<221> misc_feature
<222> (1)...(572)
<223> n = a,t,c or g

<400> 658
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gatgacctgc ctgcagagat gagccaccca ctctagggcc tcctgtctgc tgagagctgc 120
acagacaaca ggacaatcag gtacagagag gagctacact ctctgttgat agctgaacac 180
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tctgagctgt cctattgctc aataaagctc ctcttcatct tgctcaccct ccacttgcct 360
gcatatctca ttcttcctgg gcacaagata agaactcagg acctgccaaa tgaggctaac 420
agagctgtaa cacaaacagg gctcagacat gctctgtatc agtccatttc atgctggtga 480
taaagacatg cctgagactg ggaagaaaaa gaggttttat agttccccat ggctggggag 540
gcctcacaat catggcggaa cgnaacgagc ag 572

<210> 659
<211> 844
<212> DNA
<213> Homo sapiens

<220>
<221> misc_feature
<222> (1)...(844)
<223> n = a,t,c or g

<400> 659
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ccttcttccc ctcacatgtg ggactttta attccatgta tattaggctg catgaagctt 180
ccccacaacc tactgatgct cttttcatta gaaacatttc ttactctgcg tttcattttg 240
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gccattaatc ccgtccagtg tatttttcat ctcctgtatt gtagttttca tctctacaat 360
cccaacttga gcctttggtt ataacttaca tgttgctcct gcactgtttg aacatgcaga 420
atggctagtg ggcagtgag ctgaggagaa gggacagagg ggaagctcgg ctgttgggtc 480
tacgggtatg atggagacca tgcagctgaa agtaaaccgt cacccсttct gcttcagtgt 540
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aagaaagaag agataaggac ttgcctccct ttcgaggggc tgggaaacct taaccctcaa 780
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gggg 844

<210> 660
<211> 772
<212> DNA
<213> Homo sapiens

<400> 660
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gctgcagctg gactgggtga gcagaggaga gagtgtgggg ctgcatcttc ctaccctgag 180
tgtccaggag ggtgacaact ctattatcaa ctgtgcttat tcaaacagcg cctcagacta 240
cttcatttgg tacaagcaag aatctggaaa aggtcctcaa ttcattatag acattcgttc 300
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tctctctctg caaattgcag ctactcaacc tggagactca gctgtctact tttgtgcaga 420
gatccctgaa cagagatgac aagatcatct ttggaaaagg gacacgactt catattctcc 480

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ttgcgtgtac cagctgagag actctaaatc cagtgaccag gctggctggc taattaccgg 660
atttggatct tcaaccaagg tgccccaagg taggattctg tgtgtaatta cagacaaact 720
gtgctaaaca tgaggccatg actttagaac acagggtgtg gctggagcac at 772

<210> 661
<211> 920
<212> DNA
<213> Homo sapiens

<400> 661
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gtcacagcct tgtagccggg agtcgctgcc gagtgggcgc tcagttttcg ggtcgtcatg 180
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gctgtggata ccaatccact ttctctgtat gtcatgcatc cattctggaa cactatagta 300
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gggtaggttt gctctctttt ccgcctgacc ccccttggaa aagctttaca cccgcgattc 720
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accgccgccc gtctgctcac tcgccctttt tatcaaccct tcagcactcg atccgtactt 840
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cttcaatccc ttctctgccc 920

<210> 662
<211> 1372
<212> DNA
<213> Homo sapiens

<400> 662
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taagacagtc ggccggagga ttgtattttc aatataatct cctcattatt cccttcttga 120
tggttggact gtgtctacaa tgtcagagca tataggcatt acatactatg ctgtaccctt 180
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gccaaataag gtattatata tgaagtgctt taaatgtatt atttactat cttgttatcc 720
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agtactttgg gaggcccagg gtggatggat ggcttgagtc taggagttca agactagcct 900
gggcaacata gcgagatccc atctctagaa aaaaaaaaga acacaaaaat tagctggacg 960
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ctaaaaaaaa aactccattg aaatataatt ttaaaacctt ttgccgggcc gg 1372

<210> 663
<211> 1192
<212> DNA
<213> Homo sapiens

<400> 663
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gccatgccct ccagcaactc ccgccccccc gcgtgcctag ccccgggggc tctctacttg 120
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<210> 664
<211> 779
<212> DNA
<213> Homo sapiens

<400> 664
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cttgtcttcc gggcattgga gcagcccttt gagagcagcc agaagaatac catcgccttg 180
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gccatctttg gatttccact ctttggtttc ttattggctg gaattgaaga ccaacttgga 480
accatctttg ggaaaagcat tgcaagagtg gagaaggtct tttgaaaaaa gcaagtgagt 540
cagaccaaga ttcgggtcat ctcaaccatc ctgttcatct tggccggctg cattgtgttt 600
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tactttgtgg tggtcactcc gcccacggtg ggctttggtg attttgtggc agggaaaacc 720
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<210> 665
<211> 418
<212> DNA
<213> Homo sapiens

<400> 665
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ttgacacatt gaaccttgat atctgactgc ctgggtcggt catgtgctgc gtcatttgca 120
gtaagcaata tgtcctactg tccatcctgc tttgtctcct ggcatctggt tcggtggatt 180
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catttaataa gcaagactcc cttgtaattc tcaccatcat ggtaagcctt acggtttcat 300
tccctgggtt gtgcacctgc caggctggga cccaggacac ttacacttag ttcctgactt 360
gccctgatgt aggccaccct gaaaatcacg aactccaact tctacacggt ggcagtga 418

<210> 666
<211> 722
<212> DNA
<213> Homo sapiens

<400> 666
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acagcactgc tcctctagca cagcatgtcc acacacacgt atcacgccag taggccagtg 120
tgtccacata tacgcgtgca gcacagcacc actagcccag tacatccaca aacaatcgtg 180
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gacagtgcac cattaagcca gtgcgtccac acacacgtga cattacacta ttaggccggc 360
tacgtccaca cactcatgca aaattgcacc actaggccag cacatccaca cacacacgta 420
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ag 722

<210> 667
<211> 780
<212> DNA
<213> Homo sapiens

<400> 667
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gagtcatggc catgctgatg ctccccctgc tgctgctggg aatcagcggc ctcctcttca 180
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ttcccaacat gatctcgcgg agaacaggcc ctatcgtgct agggaataat atgcgaggga 660
aggtcggaac tccgaccgat ctaattcgcg tgcttcaaac acggatgcct gggcttttg 720
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<210> 668
<211> 781
<212> DNA
<213> Homo sapiens

<400> 668
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tttagaaggt atgttcatgt tcattgaggt tacatgtagg cattatagca cttgtggcat 180
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c 781

<210> 669
<211> 869
<212> DNA
<213> Homo sapiens

<400> 669
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ggccaaacat gggcgagaag ggctctgttg ctcggcatcc tgtgggccac tgcacatctg 180
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gatggaggca tcataatcta tttcctaatt atcgtttaca tgttcatggc catatctatt 360
gtctgtgatg aatacttcct accctccctg gaaatcatca gtgaatacat aggcaataag 420
aaagaaatgc aagttttaat tccaggcaga attgtttcta aattgaaaaa attaggattc 480
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aggggcttac tgcttttgaa aaaaggaagg aaagtttggg ccccgctttg cacctagcca 840
acccaatctt ataaaaaaac ccgctctgc 869

<210> 670
<211> 394
<212> DNA
<213> Homo sapiens

<220>
<221> misc_feature
<222> (1)...(394)
<223> n = a,t,c or g

<400> 670
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cttgccaatg gaccgatccc acctcattac tggaataaga aggtccccct caccccttcct 180
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<210> 671
<211> 1121
<212> DNA
<213> Homo sapiens

<400> 671
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caaaaataat aataacaaaa tattagcttt attgatgaat acctcataca ccataaaagc 180
tagtgtttat agtatagtca cagagctgca cagccatcac cacaatgtaa ttttagaata 240
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aaaacataag tccataaatg tcatttctaa gattatttta cataaatact caaatttgtt 780
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<210> 672
<211> 1245
<212> DNA
<213> Homo sapiens

<400> 672
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tgaggtcgtg caactgcttc gacccgtctc atattctcgt ccatatacac tgctgctgga 180
cacagctaat cggcattatc actatctcta cttctatcat aacaacggtt accgccgtgt 240
tcgcactctt cggcacgagt cgcctcaatg gccgtctcaa aaccctgtac actgggctca 300

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tgatatcagc aaattggaac agcagagcaa ggtgcaaaac acaggacatg gaaaaccaag     600
agaaaagtcc ataatagacg agaaattctt ccaactctct gaaatggagg cttatttaga     660
aaacagagaa aaagaagagg aacgaaaaga tgataatgat gatgagtcag ttaaaagttc     720
cagaaatgtg aacaacaaag atttttttga tccagttgaa agtgatgaag acatagcaag     780
tgatcatgat gatgagctgg gttcaaacaa gatgatgaaa ttgctgaaga agaagcagaa     840
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gtttcttttc ccttaacttt ttattttaaa aacttgcaaa cacagaaaag ttgataaaat    1140
catacagtga acatctgtat tctattcaac tggattcact agttcacatt ttgtcatatt    1200
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<210> 673
<211> 714
<212> DNA
<213> Homo sapiens

<400> 673

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ccacaaccct ggaatcctct tggctccttc agtgttggat cttttgtttc ctggatccca     180
tatcttcatt ttttcccttt ttcttagttt atgtccttgt tttggtgaca ctatactagt     240
ggctccctca gacaaggtat ataaagatac atttataata aaaatatatc catattgcat     300
atttgagaat ttcttcacat ttttatttac ttgattgttt atgttattgg agttgaaaat     360
tattttcact tagaattttg ctcagttttc ttctattctt gagagtttct gttgaagtgc     420
tttggcattc tgattcccag tcgtttacac atggcctatt ttttctgtgg aaatatttaa     480
gattttctct ttatttctga tctaagtttt tatagtgatg tgtgttgctt tgactttgat     540
tattattttt atttagttag tttttgagat agggtctcgc cctgtcacct agacaggagt     600
gcggtgacac aattatagct cagtgcaacc tcaaattcct gggctcaagc tatcctccca     660
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<210> 674
<211> 1138
<212> DNA
<213> Homo sapiens

<400> 674

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aaagttatat gagtcttggt tctataaacc attttctgtt ttttatacaa ctacttgtct     180
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tgtgtttatt attcttaatg ctatcaagtt atcatttaat atgaggtata ttttttattt     300
tgcttactta tattcagtca gaattaatga tggaatcttc ccccaccacc tccctacccc     360
aatactccag taacttatta atttattaca aagaatgacc aaaatgactt aaataagtag     420
ttatctcctg agcgtccttg acctttcttt atagtttaat tgtggtccct tgaaccagag     480
ggtgatctgc aggcattttc tttgttatca gaatgtgtga aactaggttt caggactgtg     540
tcagagaact ttttaatcat gatgcacttt ttgtcacaag aaatacttcc tcgtggaata     600
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```

caggctggcg tgcagtgtgg tgcagtctcg aatcactgca acctccaact cccggttcaa 720
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ggacctccga cctcatggga acaccctgcc tcgcctccca caattacgaa ccacagttgt 900
acccccgcc ctggaacaaa ggaacctctt ctttttatcc cccccaccgt tccgcacttt 960
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<210> 675
<211> 897
<212> DNA
<213> Homo sapiens

<400> 675
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cgggggttgg aacacgtggg gcccaagcct ttccctcccc ctgctcttat tgggtgcagt 180
tgccatggcg ctgggtgtca ggcccccagg acaggttggc ctcagcccca tcgctacggc 240
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cggcccgccc ctgtccccct ccccaccgga ctgcctctct ttgcctcgcc tcacacccct 480
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tgttcgcccg tgccccccacc ccacttaatt cttcatccga ccctcgtaca cggccgctcg 840
cgccactcct ccccgtccgc tcctctgtct ctacgaacac tcgccccggc accccccg 897

<210> 676
<211> 609
<212> DNA
<213> Homo sapiens

<400> 676
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cctgggtccc gtctgggcag cgctcctgct ctttctcctg atgtgtgaga tccctatggt 180
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agagccagag tgtgatgctg gacctggcct acggggaccg cgtctgggtg cggctcttca 420
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gccacctcat caaggccgag gacgactgag ggcctctggg ccaccctccc ggctggagag 540
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taaaatggt 609

<210> 677
<211> 999

<212> DNA
<213> Homo sapiens

<400> 677
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tggggctcta tgaggaggac agctttcatc ctgggctctg gacttctctc atttgtggcc 180
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gaacctgtgg atcctgtgaa actgcgccag tctatccgca cagttctttt caaccagtgc 480
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tgggttccca cttgtcctcc atcaccatgt ggttttcctc tggccctcat catcaccacc 840
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catctcaagt tcaaccacgg ctatggggtg tcgagcgagt ttcacgaact tctcggtaat 960
cacacggagg acgagtcatc ctggattctg agatacacg 999

<210> 678
<211> 603
<212> DNA
<213> Homo sapiens

<400> 678
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tagctgggat tacaagcacc caaccacgcc cagctaattt ttgtattttc ggtagagacg 180
ggatttcacc atgttggcca ggctagtctc gaactcatga cctcaagtga tccgcccact 240
tcggtctccc aaagtgctgg ggattacagg catgagccac ggcgccttgg ggccccaaat 300
gctcttgaaa ccggaaaccc cagggatggg agatgctcac tgagctgctg cttttatgtg 360
tgctggtgct atgtgtgttc atgtcccgcg gcagctgtct ttttgctact ataagggaat 420
tctggccacc ctgggtgggg tgtggtcggg gtgagaaccc aagcgttgga actgtagacc 480
cgtcctgtcg actgtgtgcc cctgggcatg tgtaagcctc agtttcctca tctgtaaggg 540
gggcaatgat gcctacctca caggggtgtt gtgaggatta aatgtaagga ggatagtggc 600
aac 603

<210> 679
<211> 374
<212> DNA
<213> Homo sapiens

<220>
<221> misc_feature
<222> (1)...(374)
<223> n = a,t,c or g

<400> 679
ncaaataact gtaaggaacc aagtatgact aagtgcagca gttaaggaga gtggcttgag 60

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tgctggcagc tgggttgtgc cctctgcaga gccatggcgg ccccagggct gcgcggcaca    240
catatgagga gctgtaggtg tgactggtgg gaatgaaatg accaaggccc agcgggcaat    300
tcctgggggt gtagccgcaa ccatcttctg tcggatcctg gaccatcgcc tcccagctcg    360
tgccgctcgt gccg                                                      374
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<210> 680
<211> 715
<212> DNA
<213> Homo sapiens

<220>
<221> misc_feature
<222> (1)...(715)
<223> n = a,t,c or g


<400> 680
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aaagatcgag aagcagctgc agaaggacaa gcaggtctac cgggccacgc accgcctgct    180
gctgctgggt gctggagaat ctggtaaaag caccattgtg aagcagatga ggatcctgca    240
tgttaatggg tttaatggag agggcggcga agaggacccg caggctgcaa ggagcaacag    300
cgatggtgag aaggcaacca aagtgcagga catcaaaaac aacctgaaag aggcgattga    360
aaccattgtg gccgccatga gcaacctggt gccccccgtg gagctggcca accccgagaa    420
ccagttcaga gtggactaca ttctgagtgt gatgaacgtg cctgactttg acttccctcc    480
cgaattctat gagcatgcca aggctctgtg ggaggatgaa ggagtgcgtg cctgcttacg    540
gaacgcttcc aacgagtacc agctgattga ctgtgcccag tacttcctgg acaagattcg    600
acgtgatcaa gcaggctgaa ctattgccaa cgntcaggac ctgcttcgct gccgtgtcct    660
gacttctgga atcttgagac cagttccagt tgacaagtca ncttcacatg tttga         715
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```
<210> 681
<211> 757
<212> DNA
<213> Homo sapiens


<400> 681
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gaatcatgtg tccaagggct caccctggag gtgcacagca caggtcagcc tggccagggg    120
cgaaggagac agtagagagg aagctcaggg ccttagggga ggccgggtgc aaacccgttc    180
tgcaccaagt gcactcggag tttgtgggta tgggtgtgta cccctgcagg tgtgcacatg    240
tgtgcttgca cgcacatatt tgtgcactcc tgtgcgtata catgtgtgct tgtgtatgca    300
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tgctgagaca ggaaaggggg tgaaagtgtt ggtgagggag cctggaagtt ttctcttccc    420
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agaagaaaga ttttctgact cagaagtgac tgccagtgta gcacaagcag tgtcccttgt    540
gactgtgatt ctacagttct ctgatcctca tgtttccttt agaggaaaga ggaaaaaagg    600
aactctgtgg tgggtattgg gagggaaaag aaaatagcct ggtggaggca ggagggagtc    660
gagtgtgagt aaggagcacc tgcagctttt ggaagtgaaa gcagagagag ggaaaggtag    720
ctaagacatc caggaggatc aaggggcagc gtgagag                             757
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<210> 682
<211> 1660
<212> DNA
<213> Homo sapiens

<220>
<221> misc_feature
<222> (1)...(1660)
<223> n = a,t,c or g

<400> 682
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ctggcctaaa acaaaataaa ttcttaatgg catttgtgga atgtgtttaa gagccaaaac    240
tgtgaaaatg taagctttat ctttcttttt tcctagatta tttaaagagg attgtagcca    300
caattcagat gaatgtttac aagccaaata atgatttaag agtgtgctca ataaaaaggc    360
cataggttta agaattaaat ggaataatat aaattactag gtcaacaaga atatttcatg    420
tatagtacac tgtctaagga atgcagagaa attttacaag aaacccaaga ctaaatactt    480
cattaagaac actggttact aagtaaatag atggctcatg taggaaaaag ctaatatatg    540
tagatgtaat gtcaactaag tgcatgtgac agaaatgaag aactaggaat aagaatccag    600
atttctggc caggcatttt taagtgctat tggtattcac tttatttcaa actgagcaaa    660
acaatacaac cttttacttt tttatacatt ttaaaatttc tctcatatta acattccttc    720
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gtttctcatt ttccttcttt tcccttgaag taaaccagta atttattaaa atattttata    840
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actgtctaag gaatgcagag aaatattctt gttgacctag taatttatat tattccattt   1440
aattcttaaa cctatggcct ttttattgag cacactctta aatcattatt tggcttgtaa   1500
acattcatct gaattgtggc tacaatcctc tttaaataat ctaggaaaaa agaaagataa   1560
agcttacatt ttcacagttt tggctcttaa acacattcca caaatgccat taagaattta   1620
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<210> 683
<211> 471
<212> DNA
<213> Homo sapiens

<400> 683
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gctgcatccg ttgctgcaga ggatgtgatt ttgcgctttt ctatgcttgg gcccactgtc    180
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catgcccact ccgcagcctg gacgctgctg ccggggcctc cagcccagca gcacagcact    300
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ggtcctgctg cccctgttgg gaggacgccg cctgtcctct ctgctttcac aacaacctct    420
tccttcgggt ctggctgtgg cgtcacctcc tccagggagc tgccccggcg c             471
```

<210> 684
<211> 478
<212> DNA
<213> Homo sapiens

<400> 684
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ctaggccccc agttccatct tccttttctt ttgggtgtag cagcgttgat tttctgcagg 120
tattttgaac atcagcagct gaggcaactg aacatgtttc tgtgctgtct tgcacccact 180
tctctttgga agcttcctat gtattactgc acaccttttc catgcctcct ctgtcctccg 240
cttcaacctt ccagagatgc tccagggtat cagtgggtcc catggaagac tgtctgaacc 300
aagacaagat aagatggaaa gcctcccgaa agacatgggt aggttcttag atgaacaatg 360
ggtttatttt attattttat tattattatt ttttttcga gacagtctcg ctctgtcgcc 420
caggctggag tgcagcggcg ctatatcagt tcacagcaag ctccgcctcc cgggctca 478

<210> 685
<211> 356
<212> DNA
<213> Homo sapiens

<400> 685
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cttaggtgca gtctccgaag agactactaa tgccttggaa acctggggtg ccttgcgtca 120
ggacatcaac ttggacattc ctagttttct attgagagaa catattgacg agctcatatg 180
tgataaaact ttagactcta aaaagattgc acacttcaga gctgagaaag agactttcag 240
cgaaaaagat acatattgct atttaaaaat ggaactctga aaattaagca tctgaagacc 300
gatgatcagg atatctacaa ggtatcaata tatgatacac aaggaaaaaa tgtgtt 356

<210> 686
<211> 923
<212> DNA
<213> Homo sapiens

<220>
<221> misc_feature
<222> (1)...(923)
<223> n = a,t,c or g

<400> 686
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aacaaaaaca aaaaagatgg atgggcaggg agtggaggct gtgggtagtg attgctgtcc 120
atgacccctg tctgtgagca cctgctctct aagctgaggg aatccctggt gtcatcccag 180
cagtggcgtg ttccatgctg ctgtaggcca ggaacatggt gcagccgaag tggacggcca 240
tccagtgatg acttggcccc agtggacagc tgcccagtga tgggacatct ggagtagatg 300
gccgtccaac aacagttcat tattgttgtg ctacgtctgg tgtttccagt ggctggaacc 360
actagagctc cgctccattg ggttggagcc attccagggt gggaatggcc accaggagac 420
gatgcctacc cttctcttct tgcaccaagt cagcacccat actcaggcga ggccctgtgt 480
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ttgtgaaggg atacgctgcc ttcttcctgg gctctcctgg tatcccactg gtactcagtc 600
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ggtncggaat tcgagaccag cctgaccaac gtggagagac cccatcttta ctaaaaaaaa 780
acaaagttgg ctggtggtgt ggtgcatgcc tggggccccc ctactcaggg gcctgaggcc 840
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ccacctggca aaagagggaa acc 923

<210> 687
<211> 528
<212> DNA
<213> Homo sapiens

<400> 687
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gttccaatcg ctcctgtttg tcttctgctt tggcccaggg aatgtagttt cacaaagcag 120
cttaacccca ttgatggtga acgggattct gggggagtca gtaactcttc ccctggagtt 180
tcctgcagga gagaaggtca acttcatcac ttggctttc aatgaaacat ctcttgcctt 240
catagtaccc catgaaacca aaagtccaga aatccacgtg actaatccga aacagggaaa 300
gcgactgaac ttcacccagt cctactccct gcaactcagc aacctgaaga tggaagacac 360
aggctcttac agagcccaga tatccacaaa gacctctgca aagctgtcca gttacactct 420
gaggatatta accctttacc ccattgttgg gaacgggatt tggggaata aaaacttttt 480
gacgactctc gcccgtggga atgtgaagct ggatggactc catgaatg 528

<210> 688
<211> 415
<212> DNA
<213> Homo sapiens

<400> 688
tttcgtgcca ccatcaccac cactgcggtt gctgctgcag ctgcggctgc tgctctccct 60
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ggaccactct ctttctcctg gccgggtggt gcctgccagg gttgccctgc cccagccggt 180
gcctttgctt taagagcacc gtccgctgca tgcacttgat gctggaccac attcctcagg 240
taccacagca gaccacagtt ctagacttga ggtttaacag aataagagaa attccaggga 300
gcgccttcaa gaaactcaag aatttgaaca cactgtacct gtataagaat gaaatccatg 360
cactagataa gcaaacattt aaaggactca tatctttgga acatctgtat attca 415

<210> 689
<211> 889
<212> DNA
<213> Homo sapiens

<400> 689
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tgtgattctt cccctcttgg ggctgctgct ctccctcccc gccggggcgg atgtgaaggc 180
tcggagctgc ggagaggtcc gccaggcgta cggtgccaag ggattcagcc tggcggacat 240
cccctaccag gagatcgcag gggaacactt aagaatctgt cctcaggaat atacatgctg 300
caccacagaa atggaagaca agttaagcca acaaagcaaa ctcgaatttg aaaaccttgt 360
ggaagagaca agccattttg tgcgcaccac ttttgtgtcc aggcataaga aatttgacga 420
atttttccga gagctcctgg agaatgcaga aaagtcacta aatgatatgt ttgtacggac 480

ctatggcatg ctgtacatgc agaattcaga agtcttccag gacctcttca cagagctgaa 540
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gctcctggaa cggatgtttc agctgataaa ccctcagtat cccttcagtg aaggcttcct 660
tggaatgtgt gagcaaatac cctgaccagc tcaagccatt tggagacgtg ccccggaaac 720
tgaagattca ggttacccgc gccttcattg ctgccaggac ctttgtccag ggctgactg 780
tgggcagaga agttgcaaac cgagtttcca aggtaattga aaacgtgctt tctttctcat 840
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<210> 690
<211> 784
<212> DNA
<213> Homo sapiens

<400> 690
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ccccgatgga ttcgccctgg gccgggctcc tctggctcct ccctacgctg tggtcctcat 120
ttcctgctcc ggcctgctgg ccttcatctt cctcctcctc acctgtctgt gctgcaaacg 180
gggcgatgtc ggcttcaagg aatttgagaa ccctgaaggg gaggactgct ccggggagta 240
cactccccct gcggaggaga cctcctcctc acagtcgctg cctgatgtct acattctccc 300
gctggctgag gtctccctgc caatgcctgc cccgcagcct tcacactcag acatgaccac 360
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tgggaaggtg atcctgggag agattttctc cgactacacc cccgcccagg tggtggtgaa 480
ggagctccga gccagcgcgg ggcccctgga gcaacgcaag ttcatctcgg aagcacagcc 540
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tgcgtttctg ctgatttatg gagttctgtc aactggggga cctgaagcgt tacctccgag 660
cccagcggcc ccccgagggc ctgtcccctg agctaccccc tcgaaacctg cggacgctgc 720
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acag 784

<210> 691
<211> 475
<212> DNA
<213> Homo sapiens

<220>
<221> misc_feature
<222> (1)...(475)
<223> n = a,t,c or g

<400> 691
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gggtcctggg cccagtctgt gctgactcag ccgccctcgg agtcggaggc ccctggccag 180
tgggtcaaca tctcctgcac tgggtctggc tccaacctcg ggcaggtttt tgatgtacaa 240
tggtaccagc taattccagg aacagccccc aagctcctca tctttaataa caatcgtcag 300
ccctctggag tccctgaccg attctctgcc tccaagtctg gaacctcagc ctccctaacc 360
atcaatgatc tccagcctga ggatgagtct gaatattact gccttgctat gacagcagcc 420
tcactggtgt cttcggaact gggaccaaag tcacctgcct gagtcagccc aaggc 475

<210> 692
<211> 1028

<212> DNA
<213> Homo sapiens

<400> 692
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gtgaggagga ccgggagctg gctctggagg ctgcggaggc gacgccggag agaacgaagc 180
ctcggctggg agcggatctt tcgaagatgg tttggctgcc ttggagattt ggagatctga 240
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ctttgtgtca gcctacatca ccgacgagaa cccagaagtt atgattccct tcaccaatgc 360
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cagggctgcc agctcgcacg agcacattgc agcccgcctc acggaggctg tgcacacgat 480
gctgtccagc cccttggaat acctccctcc ctgggatccc aaggactaca gtgcccgctg 540
gaatgaaatt tttggaaaca acttgggtgc cttggcaatg ttctgtgtgc tgtatcctga 600
gaacattgaa gcccgagaca tggccaaaga ctacatggag aggatggcag cgcagcctag 660
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tgaagtgatt gccaatgcct cagggtatat gtttgtaacc ttaatactag gcgcggatgg 840
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ctagtcctga tgaatcaagg atatcttcaa gaagcctact tatggaccaa acaagttctg 960
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ctgtttgc 1028

<210> 693
<211> 620
<212> DNA
<213> Homo sapiens

<400> 693
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taagatacca ctcctgctcc aaagattaca gatcccttgt cattctgact cctgggctta 120
ccctacaccc cagagatgga gcaactacta ggaataaaac ttggctgcct gtttgccctg 180
ttggctctca ctctgggctg tggccttact cccatctgct tcaaatggtt ccagattgat 240
gcagccagag gtcatcaccg gctagtcctc agactcctgg gctgtatttc tgctggagtt 300
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ctttttggag tcgctggcat tgcagtgctg tcctggggct gctggaggat cgacagtgca 540
ggacgaagaa tggggtgggg ctcatatctt cgaactccac agccatggac atttaccctc 600
accctcaaag ggtcccctcc 620

<210> 694
<211> 851
<212> DNA
<213> Homo sapiens

<220>
<221> misc_feature
<222> (1)...(851)
<223> n = a,t,c or g

<400> 694

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gaggacaatg accagactct gcttcctgtt attcttctct gtggccacca gtgggtgcag    120
tgcagcagca gcctcttctc ttgagatgct ctcgagggaa ttcgaaacct gtgccttctc    180
cttttcttcc ctgcctagaa gctgcaaaga aatcaaggaa cgctgccata gtgcaggtga    240
tggcctgtat ttctccgcca ccaagaatgg tgttgtctac cagaccttct gtgacatgac    300
ttctgggggt ggcggctgga ccctggtggc cagcgtgcac gagaatgaca tgcatggaa     360
gtgcacggtg ggtgatcgct ggtccagtca gcagggcaac aaagcagact acccagaggg    420
ggatggcaac tggggccaact acaacacctt tggatctgca gaggcggcca cgagcgatga   480
ctacaagaac cctggctact acgacatcca ggccaaggac ctgggcatct ggcatgtgcc    540
caacaagtcc cccatgcagc attggagaaa cagcgccctg ctgaggtacc gcaccaacac    600
tggcttcctc cagagactgg gacataatct gtttggcatc taccagaaat acccagtgaa    660
atacagatca gggaaatgtt ggaatgacaa tggcccagcc ataccctggg tctatgactt    720
tggggaagct taagaagact ggctcttatt actcaccgga tggtcaacgg gaatttggtc    780
cagggatccc tcaaattccc ngggttaata ccggaaagac aggccacccc ctttgtgctt    840
ggaataaagt t                                                         851
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<210> 695
<211> 995
<212> DNA
<213> Homo sapiens

<400> 695

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gtacatgcgt gcaattctcg ggtcgacgat ttcgtcttcg ctgtagacga tttcgtcgct     60
tggagtggaa gagtgggtgt ggagggcga ggctatcacg aaaagagagg aggaatcagt     120
aggaagttgc tgcctgtcct ggacccatct ggggattact actactggtg gctgaacaca    180
atggtcttcc cagtcatgta taacctcatc atcctcgtgt gcagagcctg cttccccgac    240
ttgcagcacg gttatctggt ggcctggttg gtgctggact acacgagtga cctgctatac    300
ctactagaca tggtggtgcg cttccacaca ggattcttgg aacagggcat cctggtggtg    360
gacaagggta ggatctcgag tcgctacgtt cgcacctgga gtttcttctt ggacctggct    420
tccctgatgc ccacagatgt ggtctacgtg cggctgggcc cgcacacacc caccctgagg    480
ctgaaccgct ttctccgcgc gccccgcctc ttcgaggcct tcgaccgcac agagacccgc    540
acagcttacc caaatgcctt ttgcattggc aagctgatgc tttacatttt tggccgcatc    600
cattggaaca actgcctata cttttcccta tcccggtacc tgggctttgg gcgtgaaccc    660
atgggtgtac cccggacccc ggcgccaacc tgggttttga ccgcccgggg gggccccgta    720
acctcttata agctttttaa ttttttccac cccctggata cctggattat acagggggc     780
gaataaaacc cggccgccca gtcccaggga aacaaaaaag aacctctctt cttgtgggg     840
ggcgactttt tctagttagc gccggtcaat ggggtttccc cccccccct ccttgggcct     900
tcccaggaga gctttgtgcc cttctcaaag cacgagagca ctgtgcgaaa tgggcgctct    960
ttctttcccc aaagaacttt gcgcccttgg gttcc                               995
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<210> 696
<211> 860
<212> DNA
<213> Homo sapiens

<220>
<221> misc_feature
<222> (1)...(860)
<223> n = a,t,c or g

<400> 696

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gggctccttg gagtcttgtg gaataccacc ctgcacatgt gtaggatgag actgcaagat   120
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actgggcaga aaataagaac agggagctgt gagctgcatg gttcccagag ctcacacagc 180
accgggaacc ttcgagttct gcccagccac aatggagaga ccttgcattg agtcaagagc 240
ccaggagggc cgtgcctgag atgcatggct aaaagagctt tttaggaaag gttactacag 300
acctaccatg accagggtga aaaaacaagc ctcagaagca tgaaggtgat ccacaagcaa 360
cttaggagtt gaaagaaaaa gagagagaga gagaggaggg aggaaggaag ggcggaagga 420
aaagaaacca gtactcttta aaggaagata acaaaatcca gacactcaac aatgtgacat 480
taaaaagttc catatccagt gaaaacagtc actggatatg ttctagattt taaaagacta 540
aaaagggctg gaggccaggt gcagtgactc acgcctgtaa tcccagcact ttgggaggct 600
gaggtgggca gatcacttga ggtncggagt tcgggaccag cctggccaat atggtgaaac 660
ctcgcctcta ctaaaagtgc aaagattaac cgggtgtggg gcacacgcct gtggcccagc 720
tactcgggag gctgaggcat gagaattgtt gaacctggga gcagatgttg agtgagccga 780
aatataccat ggattcagcc tggcgacgag cgagatgttc aaaaaaaaaa agaaaaaaaa 840
aagacgccgg gggtgccgcg 860

<210> 697
<211> 966
<212> DNA
<213> Homo sapiens

<400> 697
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ttttgccccc ctcttgctca tctaccttct ggtgaggatg ttcttttccg catatggctt 120
ttttatcccc ttggaacagt cctttgctag ttaatggaat atttaatgag acatttggga 180
gggaaagata gcccttgcct agtccagcct taggcaattt gggggatggg tgattacaga 240
aatgtcaggc tcttgggcag tttttccttt atctctgtca caatcagtag agtaattttt 300
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ctattccagt cccaacaagc agccagtata ctgcagtgca tactatatca tgtttcttgg 480
aagctcctgt cagctggata ataggcaatt agaagagaaa gtggacggcg ggatttaaat 540
agatcataac tggacatctg gaaaacgggg agtttgtgat gaaattaccc tgctaatgcc 600
aggttcttgc aaactttgaa aaacattata ttctaaacct catttactgt ttgggtaaaa 660
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gagtcttgct ctgttgccca ggctggagtg cagcggcatg atctcgactc actgcagcct 780
ccgcctcctg ggttcaagtg gttctcctgc ctcagcctcc ctagtagctg ggattacagg 840
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gccaggctcg tatcaaaccc ttgaccccag gtgatctgcc tgcctcagcc tcccaaagtt 960
ctggga 966

<210> 698
<211> 531
<212> DNA
<213> Homo sapiens

<400> 698
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gcatggagaa aatgttggag tgtgcattca tagtcttgtg gcttcagctt ggctggttga 120
gtggagaaga ccaggtgacg cagagtcccg aggccctgag actccaggag ggagagagta 180
gcagtctcaa ctgcagttac acagtcagcg gtttaagagg gctgttctgg tataggcaag 240
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aagaaaggct aaaagccaca ttaacaaaga aggaaagctt tctgcacatc acagccccta 360
aacctgaaga ctcagccact tatctctgtg ctgtgcaggc gcaattccat tcaggaggag 420
gtgctgacgg actcaccttt ggcaaaggca ccaggctgaa ggttttagcc ctatatccag 480
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<210> 699
<211> 559
<212> DNA
<213> Homo sapiens

<220>
<221> misc_feature
<222> (1)...(559)
<223> n = a,t,c or g

<400> 699
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agtccggaag ctaccgagcg agtccggaag ttgccgaaag ggagcagcgg ggaaggagga    120
tggcggatat catcgcaaga ctccgggagg acgggatcca aaaacgtgtg atacaggaag    180
gccgaggaga gctcccggac tttcaagatg ggaccaaggt tcgtgtctac cctgcccttc    240
tccccctctg cggcgtggtg cgcatgcgag gcgggaggag gccttaggcg agaggttgcg    300
catgcccaga gggcagcgtc cactgcccct accgctcaca tgcagaactc gacgctgatt    360
gggctgaatt taagtagggg gtgaattcgg gcctgtctgc cccgccccct ggctcggcct    420
tgtagcagca ttggtggggg aggccgtcag tcatcacaag cgggttgggg tttggggttg    480
atctcagtgc ttgngcagac cccacgctgg aggaaaccca gggccgggag tggtcctcgg    540
gtatctgggt ttcaaggct                                                 559
```

<210> 700
<211> 473
<212> DNA
<213> Homo sapiens

<400> 700
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gtgtggtgga attcctcggc tctcgccagc ccggcgcccc ggtgctgagg aatcattgac     60
atagagtaac tccacagcat gtgtcttcaa gagcttccct aaaagattaa aggttataca    120
aaacttaaaa gaagcagcaa ttctattcgc ttgttattgg acttgaaact ccctttgacc    180
tcggaaactg aagatgaggt tgccatggga actgctggta ctgcaatcat tcattttgtg    240
ccttgcagat gattccacac tgcatggccc gatttttatt caagaaccaa gtcctgtaat    300
gttccctttg gattctgagg agaaaaaagc gaagctcaat tgtgaagata aaggagatcc    360
aaaacctcat atcaggtgga agttaaatgg agcagatgct gacactggta tggagttcct    420
gctacagcgc tgttgaaagg agcttgttga tcaataaccc caataaaacc caa           473
```

<210> 701
<211> 1491
<212> DNA
<213> Homo sapiens

<400> 701
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attctgctac tcttaaaaac tagtgacgct catacaaatc aacagaaaga gcttctgaag    180
gaagacttta aagctgcttc tgccacgtgc tgctgggtct cagtcctcca cttcccgtgt    240
cctctggaag ttgtcaggag caatgttgcg cttgtacgtg ttggtaatgg gagtttctgc    300
cttcaccctt cagcctgcgg cacacacagg ggctgccaga agctgccggt ttcgtgggag    360
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gcattacaag cgggagttca ggctggaagg ggagcctgta gccctgaggt gcccccaggt 420
gccctactgg ttgtgggcct ctgtcagccc ccgcatcaac ctgacatggc ataaaaatga 480
ctctgctagg acggtcccag gagaagaaga gacacggatg tgggcccagg acggtgctct 540
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ttcttactgt gacaaaatgt ccattgagct cagagttttt gagaatacag atgctttcct 660
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tcctgtcaca agagaggatt tgcacatgga ttttaaatgt gttgtccata ataccctgag 1260
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caaacacaga actggaaaag cagatggtct gactgtgcta tggcctcatc atcaagactt 1440
tcaatcctat cccaagtgaa ataaatggaa tgaaataatt caaaaaaaaa a 1491

<210> 702
<211> 1127
<212> DNA
<213> Homo sapiens

<400> 702
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atgggggtcc ccagagtcat tctgctctgc ctctttgggg ctgcgctctg cctgacaggg 120
tcccaagccc tgcagtgcta cagctttgag cacacctact ttggcccctt tgacctcagg 180
gccatgaagc tgcccagcat ctcctgtcct catgagtgct ttgaggctat cctgtctctg 240
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gcgggccaga cgcaatcgaa cgcggacgcg ctgccgccag actactcggt ggtgcgcggc 360
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caagcacccg acccgccgac gctcagcggg ctcgagtgct acgcctgtat cggggtccac 480
caggatgact gcgctatcgg caggtcccga cgagtccagt gtcaccagga ccagaccgcc 540
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acctgccacc gggccctcct gcaccacctg atgggcacca ccagcccctg gacagccatc 660
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<210> 703
<211> 785
<212> DNA
<213> Homo sapiens

<400> 703
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aaagaataca tggacgagct tgcagctgaa gcccggggaa gtcatctaca agtgccccaa 720
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aatct 785

<210> 704
<211> 1030
<212> DNA
<213> Homo sapiens

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gatctggtaa tgaaactgga atcctcagtg aagatgccct cttgagaatc tctatcccac 420
tagactcaaa tctgaggcca gagaagtgtc gtcgctttgt ccatccccag tggcagcttc 480
ttcacctgaa tggactatc cacagcacaa gtgaggcaga cacagaaccc tgtgtggatg 540
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gaaattcaga 1030

<210> 705
<211> 1064
<212> DNA
<213> Homo sapiens

<400> 705
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acagtgtttg cagaaattac acgaggacct taaaggatac aatatagagg cagaaggcct 540
tttttcaaa gcttttttca aggagcaaac ctccaagggg cctgtatgtt tatggagatg 600
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<210> 706
<211> 413
<212> DNA
<213> Homo sapiens

<400> 706
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gtcccgggcc ggccgcggag agccccgctt catagcagag ggctacgtgg acgacaccca 180
gttcgtgagg ttcgacagcg acgccgcgag tccgaagacg gaccccgggc gccatggata 240
gagcaggaag ggccggagta ttgtgaccgc aacacacaga tcttcaagac caacacacac 300
acttaccgag agagcctgcg gaacctgcgc agctactaca accagagcga ggccggctct 360
cacattatcc agacgatgta tggctgcgaa ctgcggcccg aaggacgcct cct 413

<210> 707
<211> 311
<212> DNA
<213> Homo sapiens

<400> 707
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aaacggacca cactgttctc gagcatttca acatcaccaa gccatcagac atgggcatat 180
tctttgagtt atatcctctg ttccaagatg tacatggcat gatatttgtt gggtttgact 240
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gctgcctttc g 311

<210> 708
<211> 1196
<212> DNA
<213> Homo sapiens

<400> 708
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ctggaaagta ggtatggcct gctatgtcag gactagttct tggaattctt tgttgttttt 120
cagtcagcct tattttcttg ggtcatgttt tgaacaatat ttatcaaatg tctgtttacc 180
agacgttgtt ccagatgctt gaacaaaatg aaatgtctgc tgtcatagag tttccagtct 240
atgtaagaca gtaaacaaat gtataatata atgctagata gtgataagtg ctaaaaagaa 300
gaagaaaata ggaaagggga aagagagttt ctgtgtgatt gtatgtgaaa gtgtccatgc 360

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tttgaacaga gatctgaaat agtcagtcac gggaacattt agggagatgt tccaggcagg     480
cattgtggac aatttatgtc acaaaaaagt cacccaagtg ttaagtcaag taacatcctg     540
tatgataact atatatacat ttttttgttt tttcttaagt gaaaaacaaa cttattaggt     600
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gtttgagacc agcctggcca acatggtgaa atcccatttc tcctactcat acaaaaatta     960
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cagagcgacc ctcttgcctc acaaaacaaa acacggcctt ttctccctca ggggggacct    1140
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<210> 709
<211> 833
<212> DNA
<213> Homo sapiens

<400> 709

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taaagaccag tccaggtttc actgtttaaa acattgttct gatcatctta ttttccttct     180
aagtgaactt agaagcaata tgtttagttc tttttaatc ttatcgatat tttatgatta     240
ttgcattaat ttgtagctaa atacatgtaa aatttttat tttagccctt cttttctatg     300
gctcttaatt tttctctcat gtctgcttat gccttcagag caatgctaaa taatagtgat     360
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aggcatactt tgtcgcccaa gctggaatgc agtggcatga tctcagctca ctgcagcctt     540
gaccatctgg gctcaaccaa ttcttctgcc tcagcctccc aactcatttt ttctttaaat     600
tatttgtaga gacaagggct cgcttacaca ggctgggctt caaactctgt cttcaaacta     660
atctcccatc tcagggtcta aaagtgccgg gaataccggg ggggactaac cattacctgg     720
ggtggaagcg gtcttttggt gggtggcaa ttacctaacg gtgggggtta ataatcttaa     780
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<210> 710
<211> 490
<212> DNA
<213> Homo sapiens

<220>
<221> misc_feature
<222> (1)...(490)
<223> n = a,t,c or g

<400> 710

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acagttatat attttttctg ggttcatatg ttgcttagaa cacttcccta tacgaaaaac     180
atgaaaattt tttttcatat tttctttcat aagtgtctat ttacatatag gttatttatt     240
actcttgcgt taattttgtg gtatagtgac atagaggagt ctacctttcc ccctctaatg     300
aggtattgtc ccaacacagt gttgcataaa tctttttcc aaatgtcagc ttttatcact     360
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tatcaattct cattgtactt gagtctgttt tagattgtct cttatattga tcttttagtt    420
tataggaaag ctgctttact tnnncnnatt tctttttctt ctttgttttc gacggaccca    480
attttaaaag                                                           490
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<210> 711
<211> 1343
<212> DNA
<213> Homo sapiens

<400> 711

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gcgctatgtg cagttgggcc cccggcacca tgggcctttg gcctgcctgc tgcagagtag   180
ccctgcctgg gcagtctcca ggcactgagc aggccatctg tggccaggct gagagaatga   240
ctggctcgct taccagcgtg catgggacaa ggagctttgg agcctcaagg ggttgttgct   300
ggcctgggct agagggaaag gtgaccatcc gtctgtcctc ctgtctttct attagcgcct   360
ccatgtgagt gatggtgcct tggttcacta gccttccccc accaccccac catgccacct   420
ggtggtcttg gggcctgtgc tgtcactcca gcccctgggg aggagaggac ccagcccgga   480
gagttggggc aagggctcca catggcccaa gggcaacaga tgctcgcagg gcagctgctg   540
ccgatgctca cgctcctgcc cccctccttc ccgctgccac accccaccct gggcccccgc   600
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gcccacctgg ggccaggcca gcccctgggg cagctctgga agagcagtgt ggaggagcac   720
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gcccgtcacc agcccttcct atctgtgggg tccaagtcag gcttcccctg cggccaccag   960
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tctcaggatt gttgggggct actgaaaaga ggaatgtgtt gaatgtcgcg tttgctgtcc  1140
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tctgggaggg tgggtcagtg cacccgaggc tcagagcatc catccacccc actggccctc  1260
cttccagata ccctctctct taattggggt tctttgcatg ttaaaatact tccacaataa  1320
ataaataatt gaacaaatta aaa                                          1343
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<210> 712
<211> 648
<212> DNA
<213> Homo sapiens

<220>
<221> misc_feature
<222> (1)...(648)
<223> n = a,t,c or g

<400> 712

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ggggtgtata tagaagctac aggacaagtg taatttttat cattgcatgg ggagcattga   180
cataatttct actgcagctg agcatttttt aatatggata ataggattct gcaagtgata   240
catttggtca gagaacttaa taaactagtc aagtgggata ggtcctgtga cagaattgtg   300
tgatacaggt caaacaggag ttgggttatg gggaaaatgc cagttgaaat atgttttgat   360
ctttggagaa acctattttt tcatttaacc tgttctttaa atccagtatg ttccagaaca   420
tacaaaaatg tttaaatgtt ccatttgtaa gaggatatca tgtattttat atcaatttaa   480
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atgcagttat cctaatcatt tttctttcat ttttaccctt tattaactct tcatttgttt 540
acaaaacaaa tccactctat gaacgcaatc tctaattatg tgntttcttt cagggatcca 600
aaccttgaaa cctttgctct agatgtcagg cttgtttttg acaactgt 648

<210> 713
<211> 393
<212> DNA
<213> Homo sapiens

<400> 713
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acctgcagtt ccttctccgg gatggacgtg ggcccaact ccctgcccca ccttgggctg 120
aagctgctgc tgctcctgct gctggtgacc ctcaggggcc aagccaacac aggctggtac 180
gggattccag ggatgcccgg cctgcccggg gcaccaggga aggatgggta cgacggactg 240
ccggggccca agggggagcc gggaatcgac gccatttccc tgatcctatg acccgaagga 300
cagtaaggaa aacccgggtt tttcggacgg aaccgtaaat atggccccat gggaacctcg 360
tgggaagcaa ccgggggcca ggccccatgg tag 393

<210> 714
<211> 615
<212> DNA
<213> Homo sapiens

<400> 714
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accttgtctc ccaaatgcaa aatgtgaaat acgcaatgga attgaagcct gctattgcaa 180
catgggattt tcaggaaatg gtgtcacaat ttgtgaagat gataatgaat gtggaaattt 240
aactcagtcc tgtggcgaaa atgctaattg cactaacaca gaaggaagtt attattgtat 300
gtgtgtacct ggcttcagat ccagcagtaa ccaagacagg tttatcacta atgatggaac 360
cgtctgtata gaaaatgtga atgcaaactg ccatttagat aatgtctgta tagctgcaaa 420
tattaataaa actttaacaa aaatcagatc cataaaagaa cctgtggctt tgctacaaga 480
agtctataga aattctgtga cagatctttc accaacagat ataaattaca tatatagaaa 540
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gcaacctttc taaac 615

<210> 715
<211> 769
<212> DNA
<213> Homo sapiens

<400> 715
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aattaaatat gcagcaagtt actttgcaag gtgtcatatg gtcagtggat ggataacaaa 180
gacgcagttc ttgcttttag gaagagggaa aatttgcatg tataaatgca taaaacagct 240
acaggtaaga aaaacagatg tgataaccac aaaacagatt aattatgaag aaattaattg 300
tcttaatcat attatgctta caactaagtt ttggtaaatc catttaaaat tttggtattg 360
tatatcagta tgcattaatt cattaattca ttccattata tttattgaga ctctaccaca 420
tttcagacat gaatatatag gcatgaataa aacaaaaatg gttgcttgaa gacatggaat 480

caacctaaat gcccatcgat gacagactga ataaagaaaa tgcagtacat atacaccatg 540
gaacactatg cagccgtaaa aaagaatgag atcatgtctt ttgtggaaac atggatgaag 600
ctagaggcta ttatccttac cagacgaatg ccggaacaga aaaccaaata ccacatgttc 660
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<210> 716
<211> 743
<212> DNA
<213> Homo sapiens

<400> 716
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ggccaatcac tttggatatt agatttttaa gattgatttt ggaatttctc atccattagc 180
aggttttcac ctttctcctt aaactcatag ttttccttga aatcatacag catatttgta 240
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ttacccagca tttttaaca ttgatcaata aggcctataa ccagatttag gctagcaaca 660
ccagaggtct gggggcaagg gtggaaattg actttacatt cttagtagct aatattccat 720
aagtgcttta tatatatatt gca 743

<210> 717
<211> 630
<212> DNA
<213> Homo sapiens

<400> 717
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actacaaagt ttacctttta ctctgtaatg tggccttgta tgtttcaata caaaaataca 180
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<210> 718
<211> 432
<212> DNA
<213> Homo sapiens

<400> 718

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aactgagctt tggataattt ttcactacag ctataaccat gactgctttg ctgtagcttt   180
agccattcct gggcttggaa caggtagagt ttagcttctt ttccaacagc tactgctatg   240
ttgttgattc tgacacacag aagagcatgg tcaccatgga actcgaagtg gccgatgcgc   300
tggccctggg cgtcctgggc cgcggtgctg ctgaagctgc cccgcagggt cttaccctgg   360
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cccgtcacga aa                                                       432
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<210> 719
<211> 878
<212> DNA
<213> Homo sapiens
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cagatggggc ttcactatgg cagccagggt ggactcgaac tcctgacctt gtgatccacc   180
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ctctattctt tgaactacaa ggcaaggtca tcctcccacc cccttatcca ttcagtgaac   300
atttactgag gcgttactct gtaaagaacc ctgagagaga ccaggctgag taagacaggc   360
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gaaaatggaa atggagacaa aaagatcatg gcgcccccag tcccatggtc atttcacatt   480
ccaatttctt cttagttgga cttttgaatt aattttattt cactttgtcc cttttttttcc   540
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ctttcatcct cccctactac ctaatccttg aaaaatacaa gcccaaaatc atttcatcac   660
cagtaattgt ctttaaattg ctccaataat ttgcaaggac catggggaaa agagaaagat   720
taaaaagccc tacgcccaga gaaccagatt gtataacaag tcgaaaatca agtttactaa   780
tcaccattca tggccttgaa cttttaataa aaccttcatt gcctggaata aatccaattt   840
ttgagaaaaa cttaattgga tttaaaaatg gcgccctt                           878
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<210> 720
<211> 446
<212> DNA
<213> Homo sapiens
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<400> 720
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cttctctgcc tttctctcac tcttcctctc tgcttccccc gcctcccacc ttttccttct   180
cttcaataca ccttccctct ccccttcag gacgcctcac atccactgcc ttgccaggga   240
aggcgtgcga ctgactcagc acatctctgc cacctccatc tgcagcccaa gctggtccgt   300
gttcttgacg ggaagatacc cgatctgatc agatgaagaa cacagagtgt ggagacatga   360
agaggctttg gtgagtccac actgtaaagg gagcaggacc atgacgtctg gccccaaggt   420
tgtcaacccc aaatgcaaga tccttc                                        446
```

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<210> 721
<211> 957
<212> DNA
<213> Homo sapiens
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<400> 721
agctctatgc catcctgttt acagcgaggc aagatgaatc attatgtctg tgcattttgt 60
tttacttatc tgtgtatata gtgtacataa aggacagacg agtcctaatt gacaacatct 120
agtctttctg gatgttaaag aggttgccag tgtatgacaa aagtagagtt agtaaactaa 180
tatattttgt acattttgtt ttacaagtcc taggaaagat tgtcttctga aaatttgatg 240
tcttctgggt tgatggagat gggaagggtt ctaggccaga atgttcacat ttggaagact 300
ctttcaaatt ataactgttg ttacatgttt gcagtttatt caagactgct gtatacatag 360
tagacaaatt aactccttac ttgaaacatc tagtctatct agatgtttag aagtgcccga 420
tgtatgttaa atgtataggt agtaaaatac cactttgtaa atatcttttt gctaaaattc 480
ataggaaatg cttttggaaa ttgaattgtg aagccacctt tgtgaacagt atagtaatgt 540
ctatacttgt tcaatagttt agaggaggta ggagggaaga aattgcaaaa ggtaatatta 600
ctagtgtgtt catacttgga cattttcaga caccattttt ctatatgttt tgggcatttt 660
gttttgctct gtatatagta tatataatgg acaaatagtc ctaatttttc aacatctagt 720
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cactttgggt tgcacttcct aggaaaactt gccttttggt aagactttg ccaggaattc 840
ctctgacctt tcttattatt accgcgcccg gccggttcac ctggatgacg acaacgatgt 900
cggctgtggt caccttgggg gcccaactgg ccccttgtca tactccttga ttgagcc 957

<210> 722
<211> 925
<212> DNA
<213> Homo sapiens

<400> 722
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gcttcggcct gtgcatcgcc ttcctggggc ccacgctgct ggacctgcgc tgtcagacgc 180
acagctcgct gccccagatc tcctgggtct tcttctcgca gcagctctgc ctcctgctgg 240
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cctcctctct ggccatctcc ctggtgtttg ccgtcatccc cttctgccgc gacgtgaagg 360
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acatgcagct ggtaaggatg taccagaagg actcggccgt cttcctccag gtgctccatt 480
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ttcccaaggc tgactcccaa ggaccgggca gggaacccga ggggcctatg ccttctggat 720
aatggccctt attaatcttt ccaaggccca tggctgggct tgaagctgct ggccccaggg 780
aacggcttgt tggaactgct cgtcccccac agggggcccc ccgcttcctg gactgggaaa 840
gaaacttgcc tttgaaaaca ccagccctct tggaagaaga agacaaacct ccctcaaaag 900
gcctatagtt tatactaacg cctac 925

<210> 723
<211> 833
<212> DNA
<213> Homo sapiens

<400> 723
aaacagcgtg gtcagggaag gcttctccgc taaaggaagt agctacagga aggcaggatg 60
tgccgggcag gggagacagc aaaggcaaca gcctgagagg ggaccctgcc tgggggtcag 120
tgtggctgag tggcctgagt gaggagcaga aaggggaggc gaggtggaaa tgtgggggc 180
cagggcctgg gcctggctgg tggccctgat ggcccagggg cctctgtctc cccccaacag 240
ccctgctcct ggacatcatg acggtggccg gcgtgcagaa gctcatcaag cggcgcggcc 300

```
cgtacgagat gagccccagc ctcctggact acctcaccat ggacatctac gccttcccgg    360
ccgggcacgc cagccgggcc gtcatggtgt ccaagttctt actcagccac ctggtgctgg    420
cggtgcccct gcgcgtgctg ctggtgctct gggccctctg cgtgggcctg tcccgcgtga    480
tgatcggccg ccaccacgtc acggacgtcc tctccggctt tgtcatcggc tacctccagt    540
tccgtatgat ggagaaggtc agcatgcagt acaaaacttg ccgaatgctt atttttgtct    600
ggcgaagagc gcgtcggccc acacatacct ttgagggcag gctggtctct aaaaaggggc    660
aagacctggc caggtggctc agcctgtaat ccaaaccttt cagaggccca gtgggagcat    720
aatttaacct ccaatttgat acaagcttgg aacatggcgt cctctttttt cagacttttg    780
aaagacacgt tatctgcctt tgctgcctct ctatgagttt ctcagggccg ccc           833
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<210> 724
<211> 575
<212> DNA
<213> Homo sapiens

<400> 724

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ttccaagccc taactgggat cctcagtcta ccttgtttcc acatcccacc cacctctcgc     60
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ttaccactct cctcttcttg gcagctgtag caggggccct ggtctatgct gaagatgcct    180
cctctgactc gacgggtgct gatcctgccc aggaagctgg gacctctaag cctaatgaag    240
agatctcagg tccagcagaa ccagcttcac ccccagagac aaccacaaca gcccaggaga    300
cttcggcggc agcagttcag gggacagcca aggtcacctc aagcaggcag gaactaaacc    360
ccctgaaatc catagtggag aaaagtatct tactaacaga acaagccctt gcaaaagcag    420
gaaaaggaat gcacggaggc gtgccaggtg gaaaacaatt catcgaaaat ggaagtgaat    480
ttgcacaaaa attactgaag aaattcagtc tattaaaacc atgggcatga gaagctgaaa    540
agaatgggat cattggactt aaagccttaa ataca                               575
```

<210> 725
<211> 867
<212> DNA
<213> Homo sapiens

<400> 725

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ctgtgtgcat aggaccctgg gaccctggga tttaagtcat atggaacttg gtcaactcct    120
ccaaaatgct cccagcgctc acaggggctg ccttggtgtt tggaaggagg tggtgccaaa    180
gcagttggtt tgctggattt tgactttctt tttttaaagt ggtatttgca aatactaccc    240
cgagggcaat ggttaatgga tttgaccttt gggtcatggg ggccagggag caacactcat    300
aggagctgtg tgtgtgagtg ctgcggtgcg gcgtcgggct gctgactggc tctgccactc    360
acctctcagg ccttaagaat actgaagatt ctcacctacg attggaggcg atggtgggag    420
tggtccttaa tactgcttta tagaaaatca tagtggaggc cacgcgccgt ggctcatgcc    480
tgtagtccca gcacttcggg aagccgagat gggcggacca cgaggtcagg agatcaagac    540
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gtggtggctg actcctgtat tcccagctac tctgaaggct gaagcaggaa aatggcgtga    660
acccaggagg cggaacttgc agtgaaccga aatcgtgcca ctggactcca acctgggcga    720
cagaaagaga ctccgcctca tataaccccc tctggcgagg aatagaaata agaacccttt    780
gcggaaacca ccagggggcc ccgtgtcgcc caggggaccc tggcctcaag ttttataaaa    840
aggttgcccc aacttttttt ttccccc                                        867
```

<210> 726
<211> 861

<212> DNA
<213> Homo sapiens

<400> 726
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tgggccacga cgctaggccg gttcctcaaa ggcgcggcct ctgtacggag cagggtacgc 120
agcgtgtgtc gccccatttg tgggggccgc ggaggagggt atgtgcgctt gcgcagtccg 180
cgcgctgagc cttgcgggag gggcagttct cttgtctagc ctgtgcgcgt gtgctagggc 240
gccgcggtac gtgggcgggg aaaggcgggt gcagtcgccc gccagaccgg cagactcggt 300
tgcacgtatt gcattcatcc tctttaggtt ccgaactgac ctccagtcag gtccatcact 360
gcatcttggt atttgctgat cctctgtcct gacttgatct tgcactcagg aaagatcttc 420
aagaattacc taattttggc ctggcgcggt ggctctcgcc tgtaatccca ccactttggg 480
aggccgaggc ggttggatca actgaggtca gaaattcgag atcagcctga ccaacatggt 540
gaaaccccgt ctctactaac aataccaaaa gtaaccgggc gtggtggctc atgccctgaa 600
ctccagctac tggggggga aattgtttga aacccgggag gggcgggttc cggaaaccac 660
catggctcta ttgcacttca tattgggcta cataaacgaa tctcccgctc gcagataccc 720
atccctagaa ttacctattt tgggcgattt tgttaataaa aagaattttt ttggtttata 780
gtccaatgag ccatcccttg gtcagaaccc ccccacacgg aatatttctg catttgtttt 840
agccaaagcc tttgtgttct t 861

<210> 727
<211> 642
<212> DNA
<213> Homo sapiens

<400> 727
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ctgctacagg tcgtttcaaa ggtactgtgg ctctaccaga ccaatttctc ccttcataca 120
cattattcat ttaacagagg acagatttc aaaagaaaaa cagttcagaa ttgcaggcac 180
acatgcgcaa accctgggtc agttgaaaga ttgatttggg aatttcaata ggcaaatttg 240
gccaatgata caaatctttg gtgggagttt gctgcccaag ctaaaacctt tatacatgtt 300
ttatgaattt gcaagtttgt gatgtctgaa atcaaatgaa ctgagagttc tgctaattgt 360
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cacctagggt ctgtcgcatg tcaacaccgg ccagggctgc cagaccccgc cggcgcgaaa 480
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gctcaggagg ctaaaaccca cacccccatc ttttacgggc ct 642

<210> 728
<211> 872
<212> DNA
<213> Homo sapiens

<400> 728
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gtgacgtaga cattctcccc tgttttcttc taaaagttgc aaggtttgga ttttcttatt 120
taggtcttta atccttctag aaattatttt taggaatgat acaagttagg aatctaattg 180
tacttgtttg cttccttgta gagttattga acgttcctgt attgttcctg tattccaggg 240
gttggcagac tttgacccat gggctaactc aactcaaaac tgcctttttt ttgtaaatta 300
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gagttgagta gttgccagag atactgagtg aactccaaag cctaaaatat gtcctatctg 420
gctctttaca gaaaaagctt gcaaacccat ggtctaaaag atagtcatga aagagtagct 480

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catatttcca acagtagcag atatagtcag tgaaaataga ggaaattaca ctaaaggttg     540
taagaaggaa ggaaaacaat cttttggaca tgtaaaaaat acaaagtttg ggccgggcgc     600
ggtggctcac acctggaatc ctagcgcttt gggaggctga ggcgggtgga tcacctgggg     660
ccaggaggtc aagatcagcc ctgcccacct gggggaaccc cggcttgtgt agaatacaaa     720
aaattaccgg gcgcgggggc aagcgcccgg aatcctagca cctaggaggt tgggcaggag     780
aactgtttga ccccggagcg aagggttgac ttcgcacaga ccccacccct gccccccgct     840
ggggccatga atggggaccc ttctcaaacc cg                                   872
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<210> 729
<211> 2563
<212> DNA
<213> Homo sapiens

<220>
<221> misc_feature
<222> (1)...(2563)
<223> n = a,t,c or g


<400> 729
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cgaagctctt tgtgttcctg gccgtgctca tctgcctggt gacctcctcc ttcatcgtct     120
ttttcctgtt tccccggtcc gtcattgtgc agcctgcagg cctcaactcc tccacagtgg     180
cctttgatga ggctgatatc tacctcaaca taacgaatat cttaaacatc tccaatggca     240
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tggggcaggt ttccaacaac cttctcctac acattggccc tttggccagt gaacagatgt     360
tttacgcagt agctaccaag atacgggatg aaaacacata caaaatctgt acctggctgg     420
aaatcaaagt ccaccatgtg cttttgcaca tccagggcac cctgacctgt tcatacctga     480
gccattcaga gcagctggtc tttcagagct atgaatatgt ggactgccga ggaaacgcat     540
ctgtgcccca ccagctgacc cctcacccac catgacctgt ctgctgtccc tgtactccag     600
gcacctgcaa ccctggtcta tatctcccac aactccctgg tgactaagga aggactacag     660
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gcacctggcc tttctgtamt gagrsmacct ggtcagbctg waggggsca tttcacccca     1080
gctccatcma gggctggcag tcccvgcytg aatkdtkgga gagagctgta agttttatct    1140
tggcttttwa aaacatggac cyygccggct tggssgcaag tdggctytac acctngtaat    1200
cccagtgctt tgggnaggcc agaagtkkgg tcggkatcaa ctatgagggm agsagttccc    1260
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gagagaggca cctggcactc tgcgggcctg cacaggaaaa acagagccaa agacagaatc    1860
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gagcagcaca tcagcaggga ctggtctaga ccctcccttt cctgttcact tagctggagc    2160
taagctccag attaacccct aggttcccac tggctcctag tagaaatagt ttctgtactt    2220
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tagcagaaca ggaaggatat ttgttcatta aaggtggctt ggtcttacag ctgggtgcag   2280
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cccgggattt caattttaac atgagcagca acattagcaa gaccccgttt aaaaaaaaaa   2400
aaagagctgg gcgtaatggc gcacacctgt aatcccacct actcaggagg ctgaggcagg   2460
agaatcactt tgaacccggg aggcagaggt tgcagttgag ccaagtttcg caccattgca   2520
ttccagcngg gggcaacaag gggcgagatt ctgttttcaa aaa                      2563
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<210> 730
<211> 988
<212> DNA
<213> Homo sapiens

<400> 730

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tgggacagaa tgagggccca aactcgggcc tgctaagcca ccagtccagg agtgattcta    180
cgacatggta ttgccccctc ataagactgt tcagcttccc agactgcact tggtgtggct    240
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tattctagcc tggatgagcc tgggtccttg ttactgctgc ctattcacca ttcctaccct    420
cctggaaatt agcctcatag tatcacttgc ctaattattt tatttaattt gcacagcaca    480
gaactaaagc acagtggtga caatggccgg gaatcaagta aagtgaggta ccctatatcc    540
catcttgctg actacccagt gtagtgcctg gaacatacaa actgcacatt catacttttt    600
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atgaaaatac tgatgcttaa tatttagcag aaccaaaaaa attgtggtat aattacaact    840
ctgtaaagac aaagtaggcc gggcgcggtg gcacacgcct gtggtcccag cactttggga    900
ggccgaggcg ggtggattgc ttgagctcag gagttcaaca ccaccctggg caacatggtg    960
aaaacttgtc tctactaaaa tacaaaaa                                        988
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<210> 731
<211> 848
<212> DNA
<213> Homo sapiens

<400> 731

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atggttaata ttaggaaatt cattcagtgt ttgtttaccc aaggcttaat atatgccgga    180
ctttgtctgg ttgtagagat actgtgagga atgagttctt gctacttgcc catagaaggc    240
attcagtcca aatgcactgc gttttagaga ttcttgtttc tgttctcggt cttactcatc    300
atcttcttct tagggacagg gatcattata ggctagtgag gctgatggga gacgtaggtg    360
gtgagggaga actgaaggca atgtggaggg tgtgtctgag tgtgtgtagg gttgataaat    420
gatgctagag aagtaagaaa aggctagatc ctgtaccaga gatgtttagg agctcagatt    480
ttatcctaag agtcatagga gaggtactga agggagaagg tcatgatcag atttgcgcag    540
taaaatgatc actctggcgc cggacgtggg ggctcactcc tgtaatccca gcactttggg    600
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gaaaccctgg cttactaata tccaaaacta gcccggcgtg tgtggctgcc tgacacccat    720
ttctccgttg gttatgcaaa caacccttga ccttgaaacc gacgttcact aattctattt    780
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ctgatcct                                                              848
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<210> 732
<211> 454
<212> DNA
<213> Homo sapiens

<400> 732
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gagtgcatga gtgagggatg ttctctggag ctgaaaaaca gtaaattgaa ggaaaagaga 180
taaagcgatt tgcagagaaa ctgtagagat ttcctaaggg ccctttcagt attaagacaa 240
ttaaaaatta tagctgttcc tccttcagga aaccagagcc ccaacctact ctttttgtta 300
tctatgctgt tgtgttcact aaggacgcta ttctgtttat attatattca gtgacttaca 360
gcctgaggtc tctatgtcgt tccatcatga ttgcctcaaa aattagtgag gtttccatca 420
gtggataatt ttttattatt aaaaatttat gaag 454

<210> 733
<211> 897
<212> DNA
<213> Homo sapiens

<400> 733
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agttgctgtt ttaaattagt tgcatccagt tcatatcaat gtctgcatgc tttctagtct 120
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tggtaaatat gtcagtggcc ttttaactaa acatcaaaat gtactttaac cagttagtct 240
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ctcttctctc gccttacccc cgccgttcca ccagacagac tctgtgatcg tgctcgtccg 840
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<210> 734
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<212> DNA
<213> Homo sapiens

<400> 734
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gaggatatgg ggtttctttc actgactttg tatttgttga cttcactaaa caaaatgctc 180
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gcggaagttt ggaatgctgt gagccatttt aaggatatga ctagattctt caaatatcag 360
aaggatacca tttccaagag ggatgagatc cattctttgt aattctagga ggacaactct 420

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cctgacagca tgcatatgtg gcaaaccaga aggattacaa tgcaggagaa gtctaaggcc 660
tagcctaaac tttacctatt gtttccagcg cttccatttc tttttcttaa tctttcatta 720
ttgaaagaga tatttcgtcg acggcgccgc gagattccaa aatttatgaa tcgtaggttc 780
ctggaaaaac cccgcaggt ttcactttaa ctgggcatgg ggggcaccat atta 834

<210> 735
<211> 724
<212> DNA
<213> Homo sapiens

<400> 735
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gcttttgaaa atgatagctt ctgtaatatt tctttattca tttatttcaa tgtttaaaac 180
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gtgaagatat atactaaata ggtcgggcac agtggcttac acctgtaatc ccagcactct 480
gagaggccga ggtggacaga ccacttgagc ccaggagttc cagatcagct tggacaacat 540
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aacccagcta cgcaggaggt tgaggcatga gaatggcttc aacctgggag atagcattga 660
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aaaa 724

<210> 736
<211> 355
<212> DNA
<213> Homo sapiens

<400> 736
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accaccagag gatggagtac agatgaggct aatacttact tcttggaatg tacctgttct 120
tggtctccat ctctgcccaa aagctgcccg gaaatcaaag accaatgtcc tagtgcattt 180
gatggcctgt attttattcg tactgagaac gctgttatcc accatacctt ctgtgtcatg 240
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gacaccccgt agaattatac ccagactctt ttaagggaaa aactgctcat tattg 355

<210> 737
<211> 228
<212> DNA
<213> Homo sapiens

<400> 737
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aaactggtac tcatgaatga tgaatgtctg aggctcacat tctggctgca ctgcaatgct 180

aaacactaca gatatagcat gctgggcttt cctaaactga catctgtt 228

<210> 738
<211> 708
<212> DNA
<213> Homo sapiens

<400> 738
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gttaaaatta tttctgctgt atttgtagaa cagaagtttt gggattttgt aaaataatga 180
ccagagacta agaattccca tgccaccccg tatcactgtg gaagatggag aagtgaggaa 240
ctgtacctgc gggtgagccc tggtgccatg ttgagtgtgg gaatcaggag agctgcagtg 300
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tgaaccccga gactgcaccc actggcctcc agcctggggc aaaaaagccg ggacttcctt 660
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<210> 739
<211> 1798
<212> DNA
<213> Homo sapiens

<400> 739
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caaaaaaact gtggccccaa gtggaaccct tgaccttttc ctcagataat ctgtgtatgt 180
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taacatgaaa aattcactct tttattttgg aaaaaaagtt aacttttcat actaacaaac 360
agaacaagat ttaaggtaaa tttcttaaac attatccaga aaaataacaa gatttatagt 420
atctacttct ggtactaata tacacaaaag gccaaaacca tgcctattct gcaggtgtag 480
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ctaagagtgg aactccatta atgcttccag gtgcatcaag gtctactcct ttgacttttg 1740
tccctgtagt tccataaact cttccccctg tcttgacgaa atcgtcgacc cgggaatt 1798

<210> 740
<211> 393
<212> DNA
<213> Homo sapiens

<400> 740
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aaaaatattc cgagcattat ctggcacagg gcgtgaggtg gtgacattga gacaagtggt 180
cgaggcaagg gtgggaatag tgaccaagcc gtctctccca ggaacccaga ttatcgtcct 240
ctctggaggc gtcatcatca cggggcagtg cgcaagaggg gagggagaac cggcacttct 300
tcatatcagt tcttcttgaa atgccggtgg gtggaacact acatgatcac tctccaggcg 360
ttgagaacga cgcccgctcg cgatctagaa cta 393

<210> 741
<211> 360
<212> DNA
<213> Homo sapiens

<400> 741
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acagagccta taaaagctgt cggtccttaa ggctgcccag cgccttgcca aaatggagct 120
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ggcggaggct gaaggcaatg caagctgcac agtcagtcta gggggtgcca atatggcaga 240
gacccacaaa gccatgatcc tgcaactcaa tcccagtgag aactgcacct ggacaataga 300
aagaccagaa aacaaaagca tcagaattat cttttgctat gtccaacttg gttccgaaag 360

<210> 742
<211> 908
<212> DNA
<213> Homo sapiens

<400> 742
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accactgcgg atctcggaag gaagaaatga tgtaaatcac tcatccaaac cttaaggtca 120
aaggtgagaa ggaaggtcag gaagaacatg gcctggccaa atgtttttca aagagggtct 180
ctgctgtccc agttcagcca tcatcatgtt gtagtgttcc tgctcacttt cttcagttat 240
tcgttgctcc atgcttcacg aaaaacattt agcaatgtca aagtcagtat ctctgagcag 300
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catttgttcc ccagtgcaga gaaagcgact cttttcctcg gcacactgga taccattttc 420
ctcttctcct atgctgtggg cctattcatc agtggcatcg ttggggatcg gttgaatttg 480
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aaagccggac gaggagttgt ttttggtctc tggagtgcct gtgcttcggt gggcaacatt 720

ttgggagcgt gcctagcttc ttctgttctt cagtatggtt atgagtatgc ctttctggtg 780
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ccattaat 908

<210> 743
<211> 434
<212> DNA
<213> Homo sapiens

<400> 743
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tcttgcgctg tgtccccatc tctaatcact tatacttcta ttggtacaga caaatcttgg 180
ggcagaaagt cgagtttctg gtttcctttt ataataatga aatctcagag aagtctgaaa 240
tattcgatga tcaattctca gttgaaaggc ctgatggatc aaatttcact ctgaagatcc 300
ggtccacaaa gctggaggac tcagccatgt acttctgtgc cagcagtgaa agggggtctg 360
gggccaacgt cctgactttc ggggccggca gcaggctgac cgtgctggag gacctgaaaa 420
acgtgttccc accc 434

<210> 744
<211> 786
<212> DNA
<213> Homo sapiens

<400> 744
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gggtccctgg tcagccccgc tccctagagc aggagatctt gagtgggaga acattcttgt 120
tgtagccaca gctgaggccc tggaccagct ctctccacac cgcatgctcc gagttgggac 180
tctaaggagt ctaggaattt tcattcaaac ttggccttac aggtcactca tcagaaaaat 240
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taaatattta gccacatggt atgtaggctt ccatgtacac tcttgccctg gcccctgaaa 360
cataagcagg gggctcttct gtacatttgc ccagcttccc tgccagcctt taaccccagg 420
aacctctcag tctacctcct cttttctgcc tctgaatccc tacctttaaa gtcagaacag 480
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cacttgacat cagtagttca agaccagcct ggccaacatg gtgaaacccc atccttacta 600
aaaatacaaa aattagccag gtgtggtggc gggcacctgt aatcccagct actcaggagg 660
ctgaggcagg agaatcactt gaacccagga ggcagagttt gcagtcagcc aagatcacgc 720
cactgtactc cagcctggat gacacagcga gactccgtct caaaataaat acaaaaaaaa 780
aaaagg 786

<210> 745
<211> 379
<212> DNA
<213> Homo sapiens

<400> 745
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gcggggacat cgtgatgacc cactctccag actccctggc tgtgtctctg ggcgagacgg 120
ccaccatcga ctgcaggtcc agccagagtg tcctctacca cgccaacaat aaaaactact 180

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taacttggta ccagcagaga ccacgacagt ctcctaaagt gctcattttc tgggcatcta    240
cccgggaaac cggtgtgcct gaccgattca ctggcagcgg gtctgggaca gattattcgc    300
tcaccataag cagcctgcag gctgaagatg tggccactta ttactgtcaa caatattatg    360
attctccgat caccttccg                                                  379


     <210> 746
     <211> 440
     <212> DNA
     <213> Homo sapiens


     <400> 746
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ggagcctaca taaaagagag taaagagggg caaaaaccca gatcagaatg caggcgacgt    120
ccaaccttct caacctcctg ctgctgtctt tgtttgccgg attaaatcct tccaagactc    180
acattaatcc taaagaaggg tggcaggtgt acagctcagc tcaggatcct gatggcggg     240
gcatttgcac agttgttgct ccagaacaaa acctgtgttc ccgggatgcc aaaagcaggc    300
aacttcgcca actactggaa aaggttcaga acatgtccca gtctattgaa gtcttaaact    360
tgagaactca gagagatttc caatatgttt taaaaatgga aacccaaatg aaagggctga    420
aggcaaaatt tcggcagatt                                                 440


     <210> 747
     <211> 942
     <212> DNA
     <213> Homo sapiens


     <400> 747
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atacttctat atgaaatatg taggctacaa aacagtatat acagtttaat accattttta    120
tggaaagaaa aataaccata tatacaaaat catgcataag aaaaaaataa tataaggatg    180
tacataccaa atattaataa taatggctat ctctggatag tggaatcaga gggattatgt    240
aattttcctg ataaattttc ctgtcctcca aacagcatcc gcttcatact attatttctt    300
ggttgtaatt agtttgatat aattctcttc agaaaggctc tgtttcacta tatatacctc    360
aaagcatact tttgatgcag cttctgcaat tcccatctaa aaagtagata acacttgctc    420
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gccaggcatg gtggctcaca cctgtaattc caaaaccttg gcaggaagat cgattgaggc    540
caggagctca agacgagcct gggcaacata gaaagaccct atctttacaa aaaaaacttt    600
aaaaattagc caggtgtaat agcacatgcc tgtctgtaat cccagctact tggcaggctg    660
gaaggtcaag gctgcagtga gccatgatca tgccactgca ctccagccta ggtgacagag    720
caagaactca tctctaaaaa aaaattttta aataaagcaa aatatgccac agcatagatc    780
tgattgtaga aaattattat atggagaact gaaaatctc ctaatcaaga caaaaatttt    840
aaatagagga aaaaaatact atctatcatt agttcaagtt tccatttaaga gtagagtgtg   900
aagtagctcc aagttcagag ctggagaatt ttgcatctct cc                       942


     <210> 748
     <211> 1050
     <212> DNA
     <213> Homo sapiens

     <220>
     <221> misc_feature
     <222> (1)...(1050)
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<223> n = a,t,c or g

<400> 748
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tcgattacgg acatgcaaga tggaaaattg gttgtggcat ccagataagg gaaaacaagt 120
aggacaccag attgtataca ctgtgatcaa aaccatgtga aaaacacatg catgaagagg 180
actgggaaga aatacacaag aagtggttgc attagggtga gaaggagtat tcatgttttt 240
ctcatccgtc tttttcaaac cttttgtaat gggtggtttt attaatttta taatggaaaa 300
tgttaattta aaagcaagtt atttacagtt tagtaagctc atggcaggga aaggctgggc 360
tctgtttatt gctcttactt tttcccaacg cctactccca tgcctggcaa ttatagagat 420
aataaatgtg ggtgtggaat gagtgcccac tgggaaacct ctcagaggac tttgacccag 480
gaacatattt gcacagggtt tccctcagct ggagaaggtt tctctgggag agcaccagcc 540
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tcccagcaag gtgtaggggc ccttctggcc atttgacatc accagggcag ttagtgctga 660
tacaaaccac agagaatgaa caaactccaa ctcaaacggg aatggatttt atgtcattct 720
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gtagcaagac ccagtctcta caaaaaaatt ttttgttctg ttttgttttt gagacagagt 900
ctcaactctg tcgtctaggc tggagtgcag tggtttgatc ttgggtnatt agtttctttt 960
tttgtgggtg ttgtgtttaa gtttttgttt tgggttaaat taatctggtc ttgggaatcc 1020
ttctttttat cgttggtgga gatttaaccg 1050

<210> 749
<211> 390
<212> DNA
<213> Homo sapiens

<220>
<221> misc_feature
<222> (1)...(390)
<223> n = a,t,c or g

<400> 749
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acagacagcc actttcacct gctctggaga tgacttgggg aacaagtata tttgttggta 180
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tgtgttcgga ggaggcaccc agttgaccgn 390

<210> 750
<211> 441
<212> DNA
<213> Homo sapiens

<400> 750
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cttgaaggga atccaactta gctttaatta acattcttaa ccttcttacc tctctggatc 120
tcagttgtct catctgtaaa aaggagataa aaattattta cctgcctgaa catgaggtgg 180
aggaccatcc tgctacagta ttgctttctc ttgattacat gtttacttac tgctcttgaa 240
gctgtgccta ttgacataga caagacaaaa gtacaaaata ttcaccctgt ggaaagtgcg 300

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aagatagaac caccagatac tggactttat tatgatgaaa tcgttttaga agagcttggt    360
ggtccatgcc tatatcttga agggaatcca acttagcttt aattaacatt cttaaccttc    420
cgcacgcgtg ggtcgacccg g                                               441
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<210> 751
<211> 449
<212> DNA
<213> Homo sapiens

<400> 751

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gtggggaatt ccccagcaat cagactcaac agacggagca actgccatcc gaggctcctg     60
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tgtgggcact ggtgcgagca caggggacag ggtctgtgtg tccctcctgt ggggctcca     180
aactggcacc ccaagcagaa cgagctctgg tgctggagct agccaagcag caaatcctgg    240
atgggttgca cctgaccagt cgtcccagaa taactcatcc tccaccccag gcagcgctga    300
ccagagccct ccggagacta cagccaggga gtgtggctcc agggaatggg gaggaggtca    360
tcagctttgc tactgtcaca gactccactt cagcctacag ctccctgctc acttttcacc    420
tgtccactcc tcggtcccac cacctgtac                                      449
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<210> 752
<211> 524
<212> DNA
<213> Homo sapiens

<400> 752

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<211> 474
<212> DNA
<213> Homo sapiens

<220>
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<223> n = a,t,c or g

<400> 753

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cccctgggcc tggctccaga cacctttgac gatacctatg tgggttgtgc agaggagatg    180
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tactgggagt tgaatcangc cgtgcggacg ggcggaggct cccgggagct ctacatgagg 420
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<210> 754
<211> 1222
<212> DNA
<213> Homo sapiens

<400> 754
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<210> 755
<211> 667
<212> DNA
<213> Homo sapiens

<220>
<221> misc_feature
<222> (1)...(667)
<223> n = a,t,c or g

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ggggaaa 667

<210> 756
<211> 411
<212> DNA
<213> Homo sapiens

<220>
<221> misc_feature
<222> (1)...(411)
<223> n = a,t,c or g

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<210> 757
<211> 388
<212> DNA
<213> Homo sapiens

<400> 757
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gctgtgggcg ctgctgagcc tatggctgtg ctgcgcgacc cccgcgcctg cattgcaatg 300
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acgatcctgc ccatgcccag aaggtttt 388

<210> 758
<211> 843
<212> DNA
<213> Homo sapiens

<400> 758
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<211> 647
<212> DNA
<213> Homo sapiens

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catgcccagc ggctgccgct gcctgcatct cgtgtgcctg ttgtgcattc tgggggctcc 180
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<210> 760
<211> 796
<212> DNA
<213> Homo sapiens

<400> 760
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<210> 761
<211> 721
<212> DNA

<213> Homo sapiens

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<210> 762
<211> 716
<212> DNA
<213> Homo sapiens

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<210> 763
<211> 642
<212> DNA
<213> Homo sapiens

<400> 763
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<210> 764
<211> 2280
<212> DNA
<213> Homo sapiens

<400> 764
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<210> 765
<211> 555
<212> DNA
<213> Homo sapiens

<400> 765

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<210> 766
<211> 2744
<212> DNA
<213> Homo sapiens

<400> 766
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<210> 767
<211> 920
<212> DNA
<213> Homo sapiens

<400> 767
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<210> 768
<211> 580
<212> DNA
<213> Homo sapiens

<400> 768
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<210> 769
<211> 531
<212> DNA

<213> Homo sapiens

<400> 769
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<210> 770
<211> 1072
<212> DNA
<213> Homo sapiens

<400> 770
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<210> 771
<211> 1271
<212> DNA
<213> Homo sapiens

<400> 771
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<210> 772
<211> 1017
<212> DNA
<213> Homo sapiens

<400> 772
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gaaaaagtta atctaactac ttacctgagg taaatttagg ttggcactgc ttcaagggaa 180
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<210> 773
<211> 980
<212> DNA
<213> Homo sapiens

<400> 773
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<210> 774
<211> 1224
<212> DNA
<213> Homo sapiens

<400> 774
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tgccaaggcc ttgctgtggt tcttaaaata atttttcaac ctaatctaat agggtgttca 180
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<210> 775
<211> 1232
<212> DNA
<213> Homo sapiens

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<210> 776
<211> 708
<212> DNA
<213> Homo sapiens

<400> 776
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<210> 777
<211> 446
<212> DNA
<213> Homo sapiens

<400> 777
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aggtccccgc agggtccggt aggccg 446

<210> 778
<211> 416
<212> DNA
<213> Homo sapiens

<220>
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<222> (1)...(416)
<223> n = a,t,c or g

<400> 778
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gngcgggagg ctgtgcgccc cgatggcagc ctggacatcc agggcatcct gccccggcac 360
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<210> 779
<211> 382
<212> DNA
<213> Homo sapiens

<400> 779
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<210> 780
<211> 437
<212> DNA
<213> Homo sapiens

<400> 780
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<210> 781
<211> 476
<212> DNA
<213> Homo sapiens

<400> 781
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<210> 782
<211> 753
<212> DNA
<213> Homo sapiens

<400> 782
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ataattttta tactcttctg catttgctaa atttcctctc attagcaggt tataccttta 180
tgatcagaaa aaaaattaaa cactgcttct aaaaaatact catctccagc acttggagat 240
cacctacctc tacattctac ccaactgagc ccaatttagt cttctcaggg ctttgcccaa 300
gaacagttca ggaatgcatg cctctgaagg ccttcctgct cttccccttc tggccttggt 360
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gttcttcctt gttggtcatc agttaatgaa gtgtattagg tgacctgagt acttgtcagt 480
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agatttggag gcgggagagc tgtactgcat cccctcaaat gttagcagtg ccaagaaatg 600
agacgctagt ctaggggggca ccacaagcag aaaggggctg tttcaaggag tcgtccgccc 660
atgggagtct cctcttctat tattcacctt gctccaagga tatcttttct tttacgtatg 720
aaaattttgt aattgttcaa ctataacacc atg 753

<210> 783
<211> 769
<212> DNA
<213> Homo sapiens

<220>
<221> misc_feature
<222> (1)...(769)
<223> n = a,t,c or g

<400> 783
tttcgtagct gatggaagat gagccccaac ttctaaaaat gtatcactac cgggattgag 60
atacaaacag catttaggaa ggtctcatct gagtagcagc ttcctgccct ccttcttgga 120
gataagtcgg gcttttggtg agacagactt tcccaaccct ctgccccgcc ggtgcccatg 180
cttctgtggc tgctgctgct gatcctgact cctggaagag aacaatcagg ggtggcccca 240
aaagctgtac ttctcctcga tcctccatgg tccacagcct tcaaaggaga aaaagtggct 300
ctcatatgca gcagcatatc acattcccta gcccagggag acacatattg gtatcacgat 360
gagaagttgt tgaaaataaa acatgacaag atccaaatta cagagcctgg aaattaccaa 420
tgtaagaccc gaggatcctc cctcagtgat gccgtgcatg tggaattttc acctgactgg 480
ctgatcctgc aggctttaca tcctgttttt gaaggagaca atgtcattct gagatgtcag 540
gggaaagaca acaaaaacac tcatcacaag gtttactaca aggatggaaa acagntttct 600
aatagttata atttagagaa gaatacagtg gattcagtct cccgggataa tagcccatat 660
tattgtgctg ggtaaaagag agtttacata cttgggattg gagaacttta aaacccccaa 720

ttatccaagt ttacgggaag gggcctatac tccggagtac cagggggg 769

<210> 784
<211> 979
<212> DNA
<213> Homo sapiens

<400> 784
cagaggctcg ggaaggggcg tggatccccg gaggcggtcc ccgggttgca gtgagggaag 60
tgggccggga ggagagatgg gcggtgaaga ggcccggccc tgccagggag gcgggcggat 120
ccgtggcagt gaccagaagg ggccggaagg gggtggccgc cggccgggcc ccgccctggg 180
gccgcctccc cgcgggttcc gttggctgtg gcggcagctg acgcttgtgg cggcggtggc 240
ttcggggtgg gcgtaagatg gcgacagcag cgcagggacc cctaagcttg ctgtggggct 300
ggctgtggag cgagcgcttc tggctacccg agaacgtgag ctgggctgat ctggaggggc 360
cggccgacgg ctacggttac ccccgcggcc ggcacatcct ctcggtgttc ccgctggcgg 420
cgggcatctt cttcgtgagg ctgctcttcg agcgatttat tgccaaaccc tgtgcactcc 480
gtattggcat cgaggacagt ggtccttatc aggcccaacc caatgccatc cttgaaaagg 540
tgttcatatc tattaccaag tatcctgata agaaaaggct ggagggcctg tcaaagcagc 600
tggattggaa tgtccgaaaa atccaatgct ggtttcgcca tcggaggaat caggacaagc 660
ccccaacgct tactaaattc tgtgaaagca tgtaagtacg caaggaggga gggagggaat 720
aaggaagacg gtgggataca actggactga agtttctgtt ttgaacatca cttctgttgt 780
taggacaaca gttaatggat atagagaact aactcagcct attataggta ggaaagaagg 840
gaactggaac actgattccc ttaagtttct tggcatgtt gccactaagc taggtgtggt 900
tctattttgt tccctttcc taaatagatt gggagtaaat ccttataact gtacttatgt 960
aagtagatgt actaacaca 979

<210> 785
<211> 550
<212> DNA
<213> Homo sapiens

<400> 785
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cggcgccgtg gcggaggagc aggagcagga gggggatgga gaggagaagg ctcctgggtg 120
gcatggcgct cctgctcctc caggcgctgc ccagcccctt gtcagccagg gctgaacccc 180
cgcaggataa ggaagcctgt gtgggtacca acaatcaaag ctacatctgt gacacaggac 240
actgctgtgg acagtctcag tgctgcaact actactatga actctggtgg ttctggctgg 300
tgtggaccat catcatcatc ctgagctgct gctgtgtttg ccaccaccgc cgagccaagc 360
accgccttca ggcccagcag cggcaacatg aaatcaacct gatcgcttac cgagaagccc 420
acaattactc agcgctgcca ttttatttca ggtttttgcc aaactattta ctacctcctt 480
atgaggaagt ggtgaaccga cctccaactc ctcccccacc atacagtgcc ttccagctac 540
agcagcaacg 550

<210> 786
<211> 932
<212> DNA
<213> Homo sapiens

<220>
<221> misc_feature
<222> (1)...(932)

<223> n = a,t,c or g

<400> 786
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ggtgtggagc gcgcggcggc tgctggtgct gctgttcacg ccgctcgcgc tgctgccggt 120
ggtcttcgcc ctcccgccca aggaaggccg ctgcttgttt gtcatcctgc tcatggcggt 180
gtactggtgc acggaggccc tgccgctctc agtgacggcg ctgctgccca tcgtcctctt 240
ccccttcatg ggcatcttgc cctccaacaa ggtctgcccc cagtacttcc tcgacaccaa 300
cttcctcttc ctcagtgggc tgatcatggc cagcgccatt gaggagtgga acctgcaccg 360
gcgaatcgcc ctcaagatcc tgatgcttgt tggagtccag ccggccaggc tcatcctggg 420
gatgatggtg accacctcgt tcttgtccat gtggctgagc aacaccgcct ccactgccat 480
gatgcttccc attgccaatg ccatcctgaa aagtctcttt ggccagaagg aggttcgaaa 540
ggacccccag ccaggagagt gaagagaaca cagggaatag aaccccaata cctntcctct 600
ctgaggaaag gctgaaactt caagctcccc ttgtgataag acttggtcag ataactgagt 660
ctggtcaatg gaatatgagt ggaaatgatg tgtgcaactt ccgggttctg tccttcctgc 720
cgggtggaat gtgaatatga tggcacctgg gacccaaaga caggagccac atcttgagag 780
atagatggca gatctgcccc tgtggctttg gatcatttac ctcagtgaac acaacaagca 840
ttatccatga aaccataggt tttgtgtgct agttctagtt tttaaaatat gaattaaatt 900
aaatacgtat ctgttaaaac ttaaaaaaaa aa 932

<210> 787
<211> 514
<212> DNA
<213> Homo sapiens

<220>
<221> misc_feature
<222> (1)...(514)
<223> n = a,t,c or g

<400> 787
tttcgtctgg agcaggcggg aaagcgctgg agagaagggg gcacctggat aaccacccat 60
cttgaaggag acctccctgc cctgcctctg ttgtccccca gagcactgcc tgatcatcct 120
ctgttcccca tcctcccagc ccttcctgct gtacctgtgg ggagctgatc tcctcagtcc 180
ccctgctttt ccccggtctg ccatcaccac cccaccacca tgcacccccct tcctggctac 240
tggtcctgtt actgtctact cctgctattc tccttgggag tccaggggtc cctgggggct 300
cccagcgctg ccccagagca agtccatctg tcttacccag gtgagccagg ctccatgact 360
gtaacttgga ccacatgggt cccaacccgc tctgaagtgc aattcgggtt gcagccgtcg 420
gggcccctgc ccctccgcgc ccagggcacc ttcgtcccct ttgtggacgg nggcattctc 480
cggcggaagc tctacataca ccgagtcacg cttc 514

<210> 788
<211> 469
<212> DNA
<213> Homo sapiens

<400> 788
cccgtaattc tcgggtcgac gatttcgtgg cgcggaggag ctctgtccgg aatcacatag 60
ataccatcgt ggaaacagca gcgcaggtca cggcgccgcg ggccctgcac cagacgctgg 120
gctctagaga ttatttctct ttattcagaa gcatacagtt gtttgctgat tgcaagaaga 180
tgtttctgtg gctgtttctg atttgtcag ccctgatttc ttcgacaaat gcagattctg 240
acatatcggt ggaaatttgc aatgtgtgtt cctgcgtgtc agttgagaat gtgctctatg 300

tcaactgtga gaaggtttca gtctacagac caaatcagct gaaaccacct tggtctaatt 360
tttatcacct caatttccaa aataattttt taaatattct gtatccaaat acattcttga 420
atttttcaca tgcagtctcc ctgcatctgg ggaataataa actgcagat 469

<210> 789
<211> 525
<212> DNA
<213> Homo sapiens

<400> 789
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aactacagct tcttggcagc gtcggtgttg gccgcgggag aaggggagac cgcggcggcc 120
cccagtgaga gcggctttcc aggacggtgc gatgtgctgc gcagcgaaga ggcaggaggc 180
cggcttcctg ggtagcggt acaggcgggc gcttactctg tgcgcttgct tccccaaccc 240
tgcaccggcc atgcgcccgg ccttggcggt gggcctggtg ttcgcaggct gctgcagtaa 300
cgtgatcttc ctagagctcc tggcccggaa gcatccagga tgtgggaaca ttgtgacatt 360
tgcacaattt ttatttattg ctgtggaagg cttcctcttt gaagctgatt tgggaaggaa 420
gccaccagct atcccaataa ggtactatgc cataatggtg accatgttct tcaccgtgag 480
cgtggtgaac aactatgccc tgaatctcaa cattgccatg ccccct 525

<210> 790
<211> 377
<212> DNA
<213> Homo sapiens

<400> 790
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tgactgtgtt gggaacatct ggagtcctta cagagataat caagttaaaa tgaggtcatt 120
agtgtgggtc ctaatccaac aactgacgcc cttatacaaa ggagaaacct ggacacagac 180
atgcacagaa gaccatgtga ccatgaaggc agagatcaga gtgatgcttc tagaagccag 240
ggaagattgc cagttaatga ccaaaagaag ccaggagaca ggcctgcaac ggattctgcc 300
tgaaggctcc cagaaggaac caaccctgac aacaccttga tcttggactt ccaacctcca 360
gagctgggag gcgacac 377

<210> 791
<211> 637
<212> DNA
<213> Homo sapiens

<400> 791
ataaacttgt tttaaattgg cttattgctg gtctctcaag gcttcctatt tttgtttgct 60
ttagtctctc taaaatttca gggaaaaact atgagtctca aaatgcttat aagcaggaac 120
aagctgattt tactactagg aatagtcttt tttgaacgag gtaaatctgc aactctttcg 180
ctcccccaaag ctcccagttg tgggcagagt ctggttaagg tacagccttg gaattatttt 240
aacattttca gtcgcattct tggaggaagc caagtggaga agggttccta tccctggcag 300
gtatctctga aacaaaggca gaagcatatt tgtggaggaa gcatcgtctc accacagtgg 360
gtgatcacgg cggctcactg cattgcaaac agaaacattg tgtctacttt gaatgttact 420
gctggagagt atgacttaag ccagacagac ccaggagagc aaactctcac tattgaaact 480
gtcatcatac atccacattt ctccaccaag aaaccaatgg actatgatat tgccctttttg 540
aagatggctg gagccttcca atttggccac tttgtggggc ccatatgtct tccagagctg 600

cgggagcaat ttgaggctgg ttttatttgt acaactg 637

<210> 792
<211> 881
<212> DNA
<213> Homo sapiens

<400> 792
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aacttcttac gctttcatga tacatttatc tagttctgtt attcaagtta aagtattata 120
cagttaagtc tatggcagag tcagattctt ttatgtgtct aactgttgcg aagtatagac 180
ttcttatatc ttatatggtg accattaaca tataacgagc atgctagcat attgttgtct 240
ttgagagcac cgtatcaact ttttgatctg tagaatgaca gaagccacat tcgatactct 300
gcgactctgg ttaataatcc tgctgtgtgc tttgcggttg gccatgatgc gtagtcacct 360
gcaagcttat ttaaatttag cccaaaaatg tgtggatcag atgaagaaag aagcggggcg 420
aataagcacg gttgagctac agaaaatggt ggctcgagtc ttttattatc tttgtgtcat 480
tgcactgcag tatgtggcgc ctctggtaat gctgcttcac acaactctgc ttttgaaaac 540
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tagtctactg tccaattctg tttactctga attaccatca gctgaaggga aaatgaagca 660
taatgcaagg caaggtccag ccgttccacc cggcatgcaa gcttatggag cagccccctt 720
tgaagatctc cagctagact tcacagagat gccaaagtgt ggagatctta ttcctagatt 780
tggactgccc ttacggatcg gctcagataa tgggctggcg tttgtggctg acttggtaca 840
gaagacggca aagtggaaag gaccccagat tgtcgttctg c 881

<210> 793
<211> 622
<212> DNA
<213> Homo sapiens

<400> 793
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ctggccgctg tgtagggctg gtgagtggct ggggctgtct gagccatgaa caacttcagg 120
gccaccatcc tcttctgggc agcggcagca tgggctaaat caggcaagcc ttcgggagag 180
atggacgaag ttggagttca aaaatgcaag aatgccttga aactacctgt cctggaagtc 240
ctacctggag gggctggga caatctgcgg aatgtggaca tgggacgagt tatggaattg 300
acttactcca actgcaggac aacagaggat ggacagtata tcatccctga tgaaatcttc 360
accattcccc agaaacagag caacctggag atgaactcag aaatcctgga atcctggca 420
aattaccaga gtagcacctc ctactccatc aacacagaac tctctctttt ttccaaagtc 480
aatggcaagt tttccactga gttccagagg atgaagaccc tccaagtgaa ggaccaagct 540
ataactaccc gagttcaggt aagaaacctc gtctacacag tcaaaatcaa cccaactta 600
gagctaagct caggttttag ga 622

<210> 794
<211> 1177
<212> DNA
<213> Homo sapiens

<400> 794
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gcagcagatg cacatggccc tgcactgaga agcgcccagc tcactgcacc tgcactcagg 120

aattgtagga ctccctctag gagttgggca catgtcgttg gtgggagccc tgtccctgcc 180
ttgagaaagc tgtaggtgtt ctgtgtccag ctgtgcacct gtcctttgtt tttgtgagtc 240
ttcttggatg cacctgaatc ctgcattcag gaggcctatc ccttgttctc tgctagcaac 300
cctgcctgct atctctcttc cggtgccctc tcagccatca gaccagagct tgcttcttcc 360
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agcacttgtg tgtacaatat ttaatttaga tcttctcagt gggcctgtgg gttagaatag 660
catgtgggat tgatgggttc atcattttac atctaaggaa aatgagcctt cggttgggac 720
ctgcctggag gcacttaaca tgccttggga ctaaacactc caaggcaaac tctgttctgg 780
caagccaaca tgccgggttc tttgtggctc aagggcgatg ggcgattcac agggccttct 840
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<210> 795
<211> 599
<212> DNA
<213> Homo sapiens

<400> 795
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tcaaattccc ttacctgagg aaggagcccg attacaagga tatttacctg ctcccacccg 120
gatctaggct ctctgtttcc tcgagtcact cccagattag tggtgtctag ctcagcactg 180
tttctgttat acttcattca taattcccag cgctgttgga cgaggatggg aagaccgcct 240
gtggccatga gccctccccg gtgctcctgg ggctaaggct ggggctgcag ccatggggct 300
gggtcagccc caggcctggt tgctgggtct gcccacagct gtggtctatg gctccctggc 360
tctcttcacc accatcctgc acaatgtctt cctgctctac tatgtggaca cctttgtctc 420
agtgtacaag atcaacaaaa tggccttctg ggtcggagag acagtgtttc tcctctggaa 480
cagcctcaat gacccccctct tcggttggct cagtgaccgg cagttcctca gctcccagcc 540
ccggtcaggc gccgggctct cctcaagggc tgtggtgctg gcccgggtgc aggccctga 599

<210> 796
<211> 709
<212> DNA
<213> Homo sapiens

<400> 796
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cattctgccc tattctgacg taggccatct atcagatggc tgactcagtc ttaactttgg 120
tgttcaccag ctgcctgctt tcagagctgt ctctggtttg ctctgatttt aggccaaccc 180
ccatctcata ccagagcagg tacggctctg gggatggctg gatcaggtgc aagtctgaag 240
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cacgagatca aaagatcgag accatccttg ctaacacggt gaaaccccat ctctactaaa 540
aatacaaaaa attaaccaag cgtgctggcg ggcgcctgta gtcccagcta cttcggaagc 600
cgagcaagaa aatgacgtga acccgggaag tggagcttgc agtgagccct aatcgcacca 660

ctgcacttca gcctgggcga cagagggaga ctccatttca aaaaaaaa 709

<210> 797
<211> 389
<212> DNA
<213> Homo sapiens

<400> 797
cgagcggaga ggagatgcac acggcactcg agtgtgagga aaaatagaaa tgaaggtaca 60
tatgcacaca aaattttgcc tcatttgttt gctgacattt atttttcatc attgcaacca 120
ttgccatgaa gaacatgacc atggccctga agcgcttcac agacagcatc gtggaatgac 180
agaattggag ccaagcaaat tttcaaagca agctgctgaa aatgaaaaaa aatactatat 240
tgaaaaactt tttgagcgtt atggtgaaaa tggaagatta tcctttttttg gtttggagaa 300
acttttaaca aacttgggcc ttggagagag aaaagtagtt gagattaatc atgaggatct 360
tggccacgat catgtttctc atttaaata 389

<210> 798
<211> 480
<212> DNA
<213> Homo sapiens

<400> 798
ccctcctgca taggctcgag acatgtagct cagcttgccc gttacctgaa cagcgggcgc 60
agtcgggccc ctgaacggtc accatgtggg ccttttcgga attgaccatg cagtccatga 120
tcaatatgat tgtctccctg ctggggttag tggccacagt caccctcatc ccggccttcc 180
ggggccattt cattgctgcg cgcctcggtg gtcagtccct cggcaaaacc agccgtcagc 240
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gggcaggggc gggaaccggg gtgcttgacc acacgtgaag actcagaact aacccaggca 360
gcctggaact cggagaggtg atgagcagaa cttactcgca ttggggaaag gatgggtagg 420
gaccctaggg tatatctggg actctggcag tggtgctttc ctccctccgc cccttgtatt 480

<210> 799
<211> 639
<212> DNA
<213> Homo sapiens

<400> 799
cggacgcgtg ggcgtatttg cgcgtatgag atgcattgtc tcttcctctg gagttgagct 60
gaatgaatac ctccgaagcc gttttgttct ccaaatggga atagctccac tataccagcc 120
tcgtcttcct tccgggggac aacgtgggtc agggcacaga gagatattta atgtcaccct 180
cttggggctt tcatgggact ccctctgcca cattttttgg aggttgggaa agttgctaga 240
ggcttcagaa ctccagccta atggatccca aactcgggag aatggctgcg tccctgctgg 300
ctgtgctgct gctgctgctg ctggagcgcg gcatgttctc ctcaccctcc ccgccccccgg 360
cgctgttaga gaaagtcttc cagtacattg acctccatca ggatgaattt gtgcagacgc 420
tgaaggagtg ggtggccatc gagagcgact ctgtccagcc tgtgcctcgc ttcagacaag 480
agctcttcag aatgatggcc gtggctgcgg acacgctgca gcgcctgggg gcccgtgtgg 540
cctcggtgga catgggtcct cagcagctgc ccgatggtca gagtcttcca atacctcccg 600
tcatcctggc cgaactgggg agcgatccca cgaaaggct 639

<210> 800
<211> 412
<212> DNA
<213> Homo sapiens

<400> 800
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gcgctgtgcg gctgctggtt ccgcctgggc ggggcccgcc cgctcatccc gttgggcccg 120
actgtggtac agacctccat gagccgatcc catgtagccc tgctgggcct gagtctgctg 180
ctcatgctcc tactgtatgc ggggctgcca agcccccctg agcaaacttc ctgcctctgg 240
ggagacccca atgtcacagt cctggctgtc tccacccctg ccaactcgcc catgttctac 300
ctggaggggt taccactcca ccttgcccac agggtggacg tgatccctct gtcctctcta 360
ggccctcttg tatctcctct ccgttgtcaa gcattgcccc ctcgcctctc cc 412

<210> 801
<211> 423
<212> DNA
<213> Homo sapiens

<400> 801
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<211> 425
<212> DNA
<213> Homo sapiens

<220>
<221> misc_feature
<222> (1)...(425)
<223> n = a,t,c or g

<400> 820
gtcctaatta gaattatgct gggcctaacc atgaccaata cgtggccata ggtggtaccg 60
gtgcgagagc gagatcagct cacttaccca ctcagactac gatccgaaag cataaccagt 120
tcagtctact ggtgccggga agactggcca aatcaggaaa tgaggaagat ctacaccact 180
gtgctgtttg ccaacatcta cctggctccc ctctccctca ttgtcatcat gtatggaagg 240
attggaattt cactcttcag ggctgcagtt cctcacacag gcaggaagaa ccaagagcag 300
tggcacgtgg tgtccaggaa gaagcagaag atcattaaga tgctcctgat tgtggccctg 360
ctttttattc tctcatggct gcccctgtgg actctaatga tgctctcaga ctacgctaaa 420
ccgan 425

<210> 821
<211> 706
<212> DNA
<213> Homo sapiens

<400> 821
ggattgagtg agcccaggag gtctaggctg cagtgagctg tgatcacacc tctgcactcc 60
agcctgggtg acagagaaag atcctgtccc aaataactaa gtaaataaga tggcctgaac 120
acttgcaccc ctaaacctgc tctgtcccag tgtgccccct cgaaaatggt ctgggttctg 180
tatgtaactg ggcctctctc ctgcagagat cctctcagac tccgaggagg accgggtatc 240
ttctaatacc aacagctatg actacggtga tgagtaccgg ccgctgttct tctaccagga 300
gaccacggct cagatcctgg tccgggccct caatcccctg gattacatga agtggagaag 360
gaaatcagca tactggaaag ccctcaaggt gttcaagctg cctgtggagt tcctgctgct 420
cctcacagtc cccgtcgtgg acccggacaa ggatgaccag aactggaaac ggcccctcaa 480
ctgtctgcat ctggttatca gcccccctggt tgtggtcctg accctgcagt cggggaccta 540
tggtgtctat gagataggcg gcctcgttcc cgtctgggtc gtggtggtga tcgcaggcac 600
agccttggct tcagtgacct tttttgccac atctgacagc cagcccccca ggcttcactg 660
gctctttgct ttcctgggct ttctgaccag cgccctgtgg atcaac 706

<210> 822
<211> 357
<212> DNA
<213> Homo sapiens

<400> 822
cggacgcggg ggcggacgct gggccttgct ccttcctcat tgggatcatc agtcagtgaa 60
ttggaaggaa atgggccatg ctggtcaaca atgttctggc ggggctgggg ggcaccctta 120
tgggcctggc caacgttgct gactcctata aaatgctcat ccttgtacga ttcctttttt 180
tcgcctactg acgcgctggg cttggagtcc cttctgggaa ctgccagcct gtggccactg 240
ctcctgagcc tcacagagct acctgccctc ctgcaaatgt gactgctgac cttctgttcc 300
gaaagacccc gctacctcta cgtaatacat aatttcgagg gacctgccag aattagt 357

<210> 823
<211> 402
<212> DNA
<213> Homo sapiens

<400> 823
cgggtcgacc cacgcgtccg atccgagcta atcagtcaat acaagtcaca tgggtttatg 60
gatatgctcc atgacaagtg gtacagggtg gttccctgtg gcaagagaag ttttgctgtc 120
acggagactt tgcaaatggg catcaaacac ttctctgggc tctttgtgct gctgtgcatt 180
ggatttggtc tgtccatttt gaccaccatt ggtgagcaca tagtatacag gctgctgcta 240
ccacgaatca aaaacaaatc caagctgcaa tactggctcc acaccagcca gagattacac 300
agagcaataa atacatcatt tatagaggaa aagcagcagc atttcaagac caaacgtgtg 360
gaaaagaggt ctaatgtggg accccgtcag cttaccgtat gg 402

<210> 824
<211> 348
<212> DNA
<213> Homo sapiens

<220>
<221> misc_feature
<222> (1)...(348)
<223> n = a,t,c or g

<400> 824
ggcacgagag aggctatgag tacaatcagg acctgatccg caagggtcag gccaacaagg 60
tgaagaaact ctccatcgtt gtctccctgg ggacagggag gnccccacaa gtgcctgtga 120
cctgtgtgga tgtgatatct agcagcatca ccggttactt acgttcgtat gtttttggtg 180
tcaattatat gtgttactct cttctttcct attgtagctc tcttcgatct ttacgccact 240
ctcgctcact gtgtgtacgc gttttctact gactctcttc tgcctgctgt gatgcttact 300
gcgcttcctc gtagtctctt cttttcgtcg tcgttgattt tatcatcg 348

<210> 825
<211> 347
<212> DNA
<213> Homo sapiens

<400> 825
ggcacgagcc ggtgggtcta cagcggaagg gagggagcga aggtaggagg cagggcttgc 60
ctcactggcc accctcccaa ccccaagagc ccagccccat ggtccccgcc gccggcgcgc 120

tgctgtgggt cctgctgctg aatctgggtc cccgggcggc gggggcccaa ggcctgaccc 180
agactccgac cgaaatgcag cgggtcatgt tacgctttgg ctgctctgtc atctgttgct 240
attgtatctc agttcgtact ggtcggtccc gggaaactgg atagtctgga gcagtcgatt 300
atgtactcgg catctctttg agttgatgga gtatcgatgt gtggttg 347

<210> 826
<211> 649
<212> DNA
<213> Homo sapiens

<400> 826
ggcacgagca cctctttgag ttccccagga agaacccatt tgcactaaaa acattattga 60
gcaaagtaga tgttactaaa gattttgaag ggatgtgtag tctttcatca cctaccttgc 120
agcactcaag tttacaaacc ctcattgggc atgtgggggt tcctgagtcc cctgtgggaa 180
gtggtttttt gccatacacc ttgtttcaga gctcagcctc agttagacag ggcaggctcc 240
agtttcctca tctacccctc tccccacagc acctctaatt aaccagccct tttcttacca 300
ctgagaaatt gaactctact aaataattac agccttgtgc cacataatga cgttttggtt 360
aacaggggac cgtgtgtata atggtggtct cataagaata taataccatg ggtttactat 420
acttttctat atttagaaat gtttagattt aagttagata tggttagatt taaaatacgt 480
aacacaggct ggacccggta gctcatgcct ggaatcccag cactttggga agccgagttg 540
ggtggatcac ctgagggcag gagtttggaa ccaccctggc caacttgggg gaccccattc 600
ttctaaaaaa cacacattac ctgggggggg gcgagccctt tatcctacc 649

<210> 827
<211> 791
<212> DNA
<213> Homo sapiens

<400> 827
ggcacgagac tgttcactac ctcctctacc tggccatggc cggcgccatc tgcagaagga 60
agagataccg gaattttgga ctctactggc tgggttcctt cgccatgagc atcctggtgt 120
tccttacagg aaacattctt ggcaaataca gctccgagat caggcctgcc ttcttcctca 180
ccatccccta cctgctggtg ccatgctggg ctggcatgaa ggtcttcagc cagccccggg 240
cgctaacccg ctgcaccgcc aacatggtgc aagaggaaca aagaaaggga ctcctgcagc 300
gtccggctga cctggccctt gtcatatatc tcatccttgc tggcttcttc actctgttcc 360
ggggcctggt ggtgcttgat tgccccacag atgcctgctt tgtctatatc taccagtatg 420
agccatacct gcgggacccct gtggcctacc ctaaggtgca gatgctgatg tacatgtttt 480
atgtcctgcc tttctgcggc ctggctgcct atgctctcac cttccctggt tgctcctggc 540
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gcttcttcgt gtgcaatctg ctgtatgcgc tgggccccca cctgctggcc taccgttgcc 720
ttcagtggcc cgcattcttc caccagccac caccctccga cccccctagcc ctccacaaga 780
agcagcattg a 791

<210> 828
<211> 348
<212> DNA
<213> Homo sapiens

<400> 828

aaaggaccat ttgcagaatt cagaaaaatt cttcagtttc ttttggctta ttccatgtcc 60
tttaaaaact tgagtatgct tttgcttctg acttggccct acatccttct gggatttctg 120
ttttgtgctt ttgtagtagt taatggtgga attgttattg gcgatcggag tagtcatgaa 180
gcctgtcttc attttcctca actattctac ttttttttcat ttactctctt tttttccttt 240
cctcatctcc tgtctcctag caaaattaag acttttcttt ccttagtttg gaaacgtaga 300
attctgtttt ttgtggttac cttagtctct gtgtttttag tttggaat 348

<210> 829
<211> 638
<212> DNA
<213> Homo sapiens

<400> 829
cccacgcgtc cgcccccaagc tggtcatgga actgatgccc atcggtctgc gggggctgat 60
gatcgcagtg atgctggcgg cgctcatgtc gtcgctgacc tccatcttca acagcagcag 120
caccctcttc actatggaca tctggaggcg gctgcgtccc cgctccggcg agcgggagct 180
cctgctggtg ggacggctgg tcatagtggc actcatcggc gtgagtgtgg cctggatccc 240
cgtcctgcag gactccaaca gcgggcaact cttcatctac atgcagtcag tgaccagctc 300
cctggcccca ccagtgactg cagtctttgt cctgggcgtc ttctggcgac gtgccaacga 360
gcaggggggcc ttctggggcc tgatagcagg gctggtggtg ggggccacga ggctggtcct 420
ggaattcctg aacccagccc caccgtgcgg agagccagac acgcggccag ccgtcctggg 480
gagcatccac tacctgcact tcgctgtcgc cctctttgca ctcagtggtg ctgttgtggt 540
ggctggaagc ctgctgaccc cacccccaca gagtgtccag attgagaacc ttacctggtg 600
gaccctggct caggatgtgc ccttgggaac taaagcag 638

<210> 830
<211> 428
<212> DNA
<213> Homo sapiens

<220>
<221> misc_feature
<222> (1)...(428)
<223> n = a,t,c or g

<400> 830
tcgatgaaga ccctgtttgt ggacagctac agtgagatgc ttttctttct gcagtcactg 60
ttcatgctgg ccaccgtggt gctgtacttc agccacctca aggagtatgt ggcttccatg 120
gtattctccc tggccttggg ctggaccaac atgctctact acacccgcgg tttccagcag 180
atgggcatct atgccgtcat gatagagaag atgatcctga gagacctgtg ccgtttcatg 240
tttgtctaca tcgtcttctt gttcgggttt tccacagcgg tggtgacgct gattgaagac 300
gggaagaatg actccctgcc gtctgagtcc acgtcgcaca ggtggcgggg tttttctnan 360
acccccctct ntcttctaca taaactgtac tccacctgcc tggaactgtc caactccacc 420
atngattg 428

<210> 831
<211> 892
<212> DNA
<213> Homo sapiens

<400> 831
cccggaagct gggaaatgac ttattaacct tcatggcctc tggtcttctg aggaagcagt 60
ctgaggagcc cgagttttga aaagggaagc aatcctccaa ggctgcgatt tccacagaaa 120
tcacatgtga gccacaggtg tcattttaaa atttctagta gcaacagaaa cgaggaataa 180
acagatggtg tttgagtcac tgaatttttg gaaggacttc aaatgtcaag cattattctc 240
catgaacagg gtgatgaggg gtctggccat caccaccacc tgcctcctga gcatgctcca 300
ggccatcacc atcagcccta gcatcttgtg gaatcatgct gctgtccagt atgtacacgg 360
tcattctctt gttcaggcat gagaggtgat accagagcct tcgcaacacc agccgctccc 420
caagagcctc cccagagaaa agggccatgc agaccagcct gtgtcttctg gaactggaac 480
acggactacc caccccctatg ttgaggcagc ttctgacagg ccttactgct tacggtcatc 540
ggtcatcagc ccacccgctt gcatctccag ctgcaagtca ctctgggccc agttctcaga 600
caaggccaag tcggccacac caggggctct ctggggagcc tggaggaagg ttgactcttt 660
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caacgggcca actggggccc ttcatagaat acccccaccc tattcttttc cgaacctctc 780
tccaaggctc tgaagactgc ctccgacgtc tgtctctcgc gcccgcgcca cccgtaaacc 840
actacgactc ttcactcatt cctgcaagtc ttcactccct ctactccgat gc 892

<210> 832
<211> 312
<212> DNA
<213> Homo sapiens

<400> 832
catagaccca tgagatgtac ttgaacggcc tgagaagatt cagtcatgca ttgttgatgg 60
gcgatatgac tgccagactt atgcggtctt tgctggctgc acaacttaca tttgtatata 120
gggtggcgca tctaatgaac gttgctcaac gcataagggg aaatcgtccc attaagaatg 180
agagactact tgcattgctt ggagataatg aaaagatgaa tttgtcagat gtggaactta 240
tcccgttgcc tttagaaccc caagtgaaaa ttagaggaat aattccggaa acagctacac 300
tgtttaaaag tg 312

<210> 833
<211> 426
<212> DNA
<213> Homo sapiens

<400> 833
gccataattt ctttcttcat tggatttgga ctaagatttg gagcaaaatg gaactttgca 60
aatgcatatg ataatcatgt ttttgtggct ggaagattaa tttactgtct taacataata 120
ttttggtatg tgcgtttgct agattttcta gctgtaaatc aacaggcagg accttatgta 180
atgatgattg gaaaaatggt ggccaatatg ttctacattg tagtgattat ggctcttgta 240
ttacttagtt ttggtgttcc cagaaaggca atactttatc ctcatgaagc accatcttgg 300
actcttgcta aagatatagt ttttcaccca tactggatga tttttggtga agtttatgca 360
tacgaaattg atgtgtgtgc aaatgattct gttatccctc aaatctgtgg tccgtcgacg 420
cggccg 426

<210> 834
<211> 445
<212> DNA
<213> Homo sapiens

<400> 834
aagcgcgcta gtagcagctc tggcagaagc aacggtggct tcgagggatg gcggcggctg 60
caacaggacc tgcagcatcc cagaggaact gactaagact ttggaacaga aaccagatga 120
tgcacaatat tatcgtcaaa gagcttattg tcacattctt cttgggaatt actgtggtgc 180
agatgctaat ttcagtgact ggattaaaag gtgtcgaagc tcagaatggc tcggaatctg 240
aggtgtttgt ggggaagtat gagaccctcg tgttttactg gccctcgctg ctgtgccttg 300
ccttcctgct gggccgcttc ctgcatatgt ttgtcaaggc tctgagggtg cacctcggct 360
gggagctcca ggtggaagaa aaatctgtcc tggaagtgca ccagggagag cacgtcaagc 420
agctcctgag gataccccgc cctca 445

<210> 835
<211> 487
<212> DNA
<213> Homo sapiens

<220>
<221> misc_feature
<222> (1)...(487)
<223> n = a,t,c or g

<400> 835
tttagatgat cccctctgaa aatgatagct gcgaanccnc cnaantnngg gtgacccacg 60
cgtgcgggat acaggcctag gctatggtaa ttgtaagcgg aagtgaaata aatattttat 120
ttgtgtgtgc atttatttaa caaacattaa ttatctcctt gattaataaa gcactgttcc 180
tgccctcaag tagttcatgg tgggctagtc caagaacaat taaatatagt atgactatac 240
atttatgtag taatctaatg tgtcatttct tgcagagaat gggaacaatt ctcctttgcc 300
caaatatgca acctcaccaa aacctaacaa cagttatatg ttcaaaaggg aacctcctga 360
gggctgtgaa agggtcaaag tctttgagga atgctcgtaa gtatcccttc caccatccgc 420
ccnnggngga aaccccccaat gggggcaaa caaggnnggg gggggcgcgg tttaaacaac 480
ccacgan 487

<210> 836
<211> 611
<212> DNA
<213> Homo sapiens

<400> 836
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ctggatgggc cttgaagcct tccacctcta cctgctcgct gtcagggtct tcaacaccta 120
cttcgggcac tacttcctga agctgagcct ggtgggctgg ggcctgcccg ccctgatggt 180
catcggcact gggagtgcca acagctacgg cctctacacc atccgtgata gggagaaccg 240
cacctctctg gagctatgct ggttccgtga agggacaacc atgtacgccc tctatatcac 300
cgtccacggc tacttcctca tcaccttcct ctttggcatg gtggtcctgg ccctggtggt 360
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gaaggtgctc accctgctgg gcctctcgag cctggtgggt gtgacatggg ggttggccat 480
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tgtcttcatc tgctgctggt tcaccatcct ttacctccca agtcagagca ccacagtctc 600
ctcttctact g 611

<210> 837
<211> 609

<212> DNA
<213> Homo sapiens

<400> 837
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actctgcaaa ggctggctga gttgtccaaa tgacagagcc cccaggggct tcgtcccatc 120
tcagacaggc attacgctgc tgccagtggc tggctggaat tccaagccag tgggttttat 180
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gccccccttcc taggggtatg taccaacatc ctgccttgcc tgagatgcca tcacctttct 300
tggggatcct aaggctggag tatgtaaagc tcctgggtct ctgtatgtgc ctgagcaccg 360
gttcttccta gactccacac agctctgtgt gttggaccca aggccctggt ggggtgggct 420
catgagggga tatcctgatc tgagggttgc aaagatccat aggagaagtg tggtttccag 480
gggtcacaca ttcactcact gcctcccttg gcttgggggg gggcctccct tggctccgtg 540
ttgctcctgg gtgggccact gccctgccct gcttttctcc attttctctg cattgaattg 600
cttccctga 609

<210> 838
<211> 11795
<212> DNA
<213> Homo sapiens

<400> 838
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ctgttgttgc tgctgtctgt tttcgtgatt cagcaagttt catctcaaga tttatcaagc 120
tgtgcaggga gatgtgggga agggtattct agagatgcca cctgcaactg tgattataac 180
tgtcaacact acatggagtg ctgccctgat ttcaagagag tctgcactgc ggagctttcc 240
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tgtaagaagt atgacaagtg ctgtcccgat tatgagagtt tctgtgcaga agtgcataat 360
cccacatcac caccatcttc aaagaaagca cctccacctt caggagcatc tcaaaccatc 420
aaatcaacaa ccaaacgttc acccaaacca ccaaacaaga agaagactaa gaaagttata 480
gaatcagagg aaataacaga agaacattct gtttctgaaa atcaagagtc ctcctcctcc 540
tcctcctctt cctcttcttc ttcaacaatt tggaaaatca agtcttccaa aaattcagct 600
gctaatagag aattacagaa gaaactcaaa gtaaaagata acaagaagaa cagaactaaa 660
aagaaaccta cccccaaacc accagttgta gatgaagctg gaagtggatt ggacaatggt 720
gacttcaagg tcacaactcc tgacacgtct accacccaac acaataaagt cagcacatct 780
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cctgccccaa ctacccccaa ggagcctgca cccaccactc ccaaggagcc tgcacccacc 1560
accaccaagg agccttcacc caccactccc aaggagcctg cacccaccac caccaagtct 1620
gcacccacca ctaccaagga gcctgcaccc accactacca agtctgcacc caccactccc 1680
aaggagcctt cacccaccac caccaaggag cctgcaccca ccactcccaa ggagcctgca 1740
cccaccaccc ccaagaagcc tgccccaact acccccaagg agcctgcacc caccactccc 1800
aaggaacctg cacccaccac caccaagaag cctgcaccca ccgctcccaa agagcctgcc 1860
ccaactaccc ccaaggagac tgcacccacc acccccaaga agctcacgcc caccaccccc 1920
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cccaccaccc ctgaggagcc cacacccacc acccctgagg agcctgctcc caccactccc   2040
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ccaactaccc ctaaggagcc tgctccaact acccctaagg agactgctcc aactacccct   2160
aaagggactg ctccaactac cctcaaggaa cctgcaccca ctactcccaa gaagcctgcc   2220
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gctccaacta cccctaaggg gactgctcca actaccccta aggagcctgc tccaactacc   2340
cctaaggagc ctgctccaac tacccctaag gggactgctc caactaccct caaggaacct   2400
gcacccacta ctcccaagaa gcctgccccc aaggagcttg cacccaccac caccaagggg   2460
cccacatcca ccacctctga caagcctgct ccaactacac ctaaggagac tgctccaact   2520
acccccaagg agcctgcacc cactaccccc aagaagcctg ctccaactac tcctgagaca   2580
cctcctccaa ccacttcaga ggtctctact ccaactacca ccaaggagcc taccactatc   2640
cacaaaagcc ctgatgaatc aactcctgag ctttctgcag aacccacacc aaaagctctt   2700
gaaaacagtc ccaaggaacc tggtgtacct acaactaaga ctcctgcagc gactaaacct   2760
gaaatgacta caacagctaa agacaagaca acagaaagag acttacgtac tacacctgaa   2820
actacaactg ctgcacctaa gatgacaaaa gagacagcaa ctacaacaga aaaaactacc   2880
gaatccaaaa taacagctac aaccacacaa gtaacatcta ccacaactca agataccaca   2940
ccattcaaaa ttactactct taaaacaact actcttgcac ccaaagtaac tacaacaaaa   3000
aagacaatta ctaccactga gattatgaac aaacctgaag aaacagctaa accaaaagac   3060
agagctacta attctaaagc gacaactcct aaacctcaaa agccaaccaa agcacccaaa   3120
aaacccactt ctaccaaaaa gccaaaaaca atgcctagag tgagaaaacc aaagacgaca   3180
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<210> 846
<211> 635
<212> DNA
<213> Homo sapiens

<400> 846
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ggctacagtt gtcgctgctt gcctggcttt gctggggagc gttgtgaggg agacatcaac 180
gagtgcctct ccaacccctg cagctctgag ggcagcctgg actgtataca gctcaccaat 240
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gtgtgtcccc agatgccctg cctgaatgga gggacttgtg ctgtggccag taacatgcct 360
gatggtttca tttgccgttg tccccсggga ttttccgggg caaggtgcca gagcagctgt 420
ggacaagtga aatgtaggaa gggggagcag tgtgtgcaca ccgcctctgg accccgctgc 480
ttctgcccca gtccccggga ctgcgagtca ggctgtgcca gtagcccctg ccagcacggg 540
ggcagctgcc accctcagcg ccagcctcct tattactcct gccagtgtgc cccaccattc 600
tcgggtagcc gctgtgaact ctcaactcac ccacc 635

<210> 847
<211> 1100
<212> DNA
<213> Homo sapiens

<400> 847
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ccagagcttg gcgatcccga agcagttggg tctgcggagg gagatgcctt cgggcagccc 180
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tttggtccac tgctttttcc ccagcccctt gcacaggacc tattacacag taggtgctca 540
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caatactcgc tgttgtatcc 1100

<210> 848
<211> 685
<212> DNA
<213> Homo sapiens

<400> 848
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ggaggcagca cagaaaaagc attctctttg tgaattgctc cgcataccca acatatgtaa 180
aagaatctgt ttcctgtcct ttgtgagatt tgcaagtacc atcccttttt ggggccttac 240
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caccctcctg gccaattgtg ttgcaccttg ggcactgaat cacatgagcc gtcgactaag 360
ccagatgctt ctcatgttcc tactggcaac ctgccttctg gccatcatat ttgtgcctca 420
agaaatgcag accctgcgtg tggttttggc aaccctgggt gtgggagctg cttctcttgg 480
cattacctgt tctactgccc aagaaaatga actaattcct tccataatca ggggaagagc 540
tactggaatc actggaaact ttgctaatat tgggggagcc ctggcttccc tcgtgatgat 600
cctaagcata tattctcgac ccctgccctg gatcatctat ggagtctttg ccatcctctc 660
tggccttgtt gtcctcctcc ttccg 685

<210> 849
<211> 413
<212> DNA
<213> Homo sapiens

<400> 849
gatttttaat aatgattcca cctgctatat tttgggtttt aattatcttc ggatggacgc 60
tcgtctacgg ttttgtatac ttcacaacgg gagaaacgat tatggacaag ttactccgtg 120
tcctctactg gattctcgtg aagaccttct tcagagagat ttcggtgtcg caccaggagc 180
gtatccccaa agataagccg gtcatgctgg tgtgtgctcc gcatgccaac cagtttgtgg 240
acggaatggt catttcaacc catctggacc gcaaggtgta ctttgtgggt gcggcctcga 300
gtttccgcaa gtacaaggtg gtgggtctct tcatgaagct gatggcgtcc atcatttcgg 360
gggagcgtca ccaggacgtg aaaaaagtgc tgaccggaat ggcgacggag aag 413

<210> 850
<211> 395
<212> DNA
<213> Homo sapiens

<400> 850
aatggatgtt ctatgtgaaa gctgagttcc ttgtttcttt ctcttgcccg tggctgactg 60
cgtgtgctct attgatgtct tgttcctggt tcttgacact gaccatcttg tctgtgaaag 120
gaggcactcc ggcgggcatg cttgatcaga agaaagggaa gtttgcttgg tttagtcact 180
ccacagaaac ccatggtaat gttcccctgt gctctgtgtg tgtaaatgcg tgtgggtgca 240
taccagactg aatgggaagg tgtctctctt gatggcttgt gccgcagtag ttctgtgtgt 300
gtgcatatat gtgtatgtat atatgttgtg tgggtgtgtg tgtttgtgaa gggatggcaa 360
cctgtccccc tcaaagccac tgccttatca tggct 395

<210> 851
<211> 904
<212> DNA
<213> Homo sapiens

<400> 851
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ataacagtgt tctcaccagt ggcgttggtg cagatgtggc caggatgtgg gagatagcca 120
tccagcatgc ccttatgccc gtcattccca agggctcctc cgtggtacaa ggaaccaact 180
tgcacagtga gtctgccagt tttctaacca gcccaaagct catcatgtgc ctaccccttg 240
cttagtaaac atgtgccctg cccttcctaa gaacagaatg aagaaagact tcttggggat 300
gacttagttt attgtagaat gtagggtgtc taaataaaag ctgctgcaca tactaagatg 360
tttagtttgt taaattatcc tattttatta tagctatttt atattaaaat ttaacaaatt 420
caggtaaaca ctatgtatta ggcaattaca gacctctaga gctattggtt ataaaagaag 480
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aaagtaacgg cgacaggggc attgacgagg gtccggggat gggcctgcaa cattatggta 900
gccc 904

<210> 852
<211> 592
<212> DNA
<213> Homo sapiens

<400> 852
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tccccaatac aactcatgag ggtttcaatg tcaccctcca caccaccctg gttgtcacga 120
cgaaactggt gctcccgacc cctggcaagc ccatcctccc cgtgcagaca ggggagcagg 180
cccagcaaga ggagcagtcc agcggcatga ccattttctt cagcctcctt gtcctagcta 240
tctgcatcat attggtgcat ttactgatcc gatacagatt acatttcttg ccagagagtg 300
ttgctgttgt ttctttaggt attctcatgg gagcagttat aaaaattata gagtttaaaa 360

aactggcgaa ttggaaggaa gaagaaatgt ttcgtccaaa catgtttttc ctcctcctgc 420
ttccccctat tatctttgag tctggatatt cattacacaa gggtaacttc tttcaaaata 480
ttggttccat caccctgttt gctgtttttg gaacggcaat ctccgctttt gtagtaggtg 540
gaggaattta ttttctgggt caggctcacg taatctctaa actcaacatg ac 592

<210> 853
<211> 436
<212> DNA
<213> Homo sapiens

<400> 853
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acactgatgt gtatacagat gctgtttccc tgctgttctc ttctaagtat gaatcccggt 120
ccccttttgca gacccagtag gtgaatccaa ttacgtagag caggggactg tggagctgtg 180
ttgtgagcag cacccaggtg atgccccatg gcagcatgtc ccacattcct tccatctttt 240
aaaaaaaatt tttctcggtg gcagtcttgc tctgtcgcct aggctggggt acagtggtgc 300
aatctcagct caccgcagcc tcaacctccc gggttcaagc aatcctccca ccttggcctc 360
ccaaagccaa agattgcagg tgtgagtcct cggctcggcg gtgggtcgac ccggaattcc 420
ggccggacga cgtcgt 436

<210> 854
<211> 266
<212> DNA
<213> Homo sapiens

<400> 854
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ttggtactga ctgtgccttt catgggatgc ttacttatcc tggtcgatgg cctaaagccc 120
aaccgtccag cttatatcca gacagggtct caagccaccc aggctggagt gcagtggcac 180
aattatggct cactgtagcc tcaccttcct gggatcaagc aatcttcttt cttcagcctc 240
cagaggagct gggaccacag atcctt 266

<210> 855
<211> 420
<212> DNA
<213> Homo sapiens

<400> 855
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caccaggagc tcaacaccct caagttccag ctgagtgctg aaatcatgga ctaccagagc 120
agacttaaga atgctggtga agagtgcaag agcctcaggg gccagcttga ggagcaaggc 180
cggcagctgc aggctgctga ggaagctgtg gagaagctga aggccaccca agcagacatg 240
ggagagaagt tgagctgcac tagcaaccat cttgcagagt gccaggcggc catgctgagg 300
aaggacaagg agggggctgc cctgcgtgaa gaccaagaaa ggacccagaa ggaactcgaa 360
aaagccacgt gtattgcgga cgaaatcgtc gacccgggaa gtccggtccg aatgctgtca 420

<210> 856
<211> 412

<212> DNA
<213> Homo sapiens

<400> 856
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tgcaagacac tggtgcattg caggtgtgtt tccgcagaag tttgatggtg acagtgccta 120
cgtggggatg agtgacggaa acccagagct cctgtcaacc agccagacct acaacggcca 180
gagcgagaac aacgaagact atgagatccc cccgataaca cctcccaacc tcccggagcc 240
atccctcctg cacctggggg accacgaagc cagctaccac tcgctgtgcc acggcctcac 300
ccccaacggt ctgctccctg cctactccta tcaggccatg gacctcccag ccatcatggt 360
gtccaacatg ctagcacagg acagccacct gctgtcgggc cagctgccca cg 412

<210> 857
<211> 403
<212> DNA
<213> Homo sapiens

<400> 857
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gcccctggct ggtttgggaa ggctgggctc ccaggctggt ggtagtggtg ggggtgattt 120
tcctcatgaa gcccccactc cgtccactac tgcctgacac ccacgaagcg agcagtttcc 180
ggagctctcc gatgtagggg cagcaggtgt agagcagctg ctggtccacc acaggcgcat 240
tgtccaagcc atgctctggg gctactgtgt ccacctcaaa ggcatatgag ggaccctctt 300
ccagaaagaa caagtcctca gggactgtgg gaatctggaa aagccagtcc agggcagcaa 360
gaagcagcag cttgttcagg aaacacatct tcccctcact ctc 403

<210> 858
<211> 439
<212> DNA
<213> Homo sapiens

<400> 858
tgagggtggc gcaggggccc cggccagccc ggggctgcag cagtgcggac agctccagaa 60
gctcatcggc atctccattg gcagcctgcg cgggctgggc accaagtgcg ctgtgtccaa 120
cgacctcacc gagcaggaga tacggaccct ggagcattgt cccaattcct tcttctaatg 180
aagaaatacg cttagttgat gatgcgtttg gaaaaatttg tcacatggtc agtgatggct 240
cttgggtggt tcgtgttcag gcagcaaaac tgttgggctc tatggagcaa gtcagttctc 300
atttcttgga gcagaccctt gacaagaagc atgtcagatc tgaggaggaa acgtactgca 360
catgagcgtg ccaaggaact ttacagttcg ggggagtttt ccagtggcag aaagtgggga 420
gatgatgctc ccaaggaag 439

<210> 859
<211> 985
<212> DNA
<213> Homo sapiens

<220>
<221> misc_feature
<222> (1)...(985)
<223> n = a,t,c or g

<400> 859

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tctgcacacg ggcatcggcc ggcatgttgg tgtggctgta cgggaccatc agatggccag     180
cactgggggc accaaggtgg tggccatggg tgtggccccc tggggtgtgg tccggaatag     240
agacaccctc atcaacccca agggctcgtt ccctgcgagg taccggtggc gcggtgaccc     300
ggaggacggg gtccagtttc ccctggacta caactactcg gccttcttcc tggtggacga     360
cggcacacac ggctgcctgg ggggcgagaa ccgcttccgc ttgcgcctgg agtcctacat     420
ctcacagcaa aacacggccg tggcagggac tggaattgac atccctggcc tgctcctcct     480
gaaagaatgt gatgagaaga tggtgacgcg aatacacaac gccagccagg ctcagctccc     540
atgtcttcct tatgattgcg ttaaggggga gctacggact tgcctagcgg gcaccccttg     600
gaataccctc ttgcccccgg gaacggtggt tttccagcct acgccccgaa ccccgagaat     660
gcatccacgc gcctcgtttt gctgaattga ngatccttgg acgtccttgc atcccacatc     720
gtggcgaaat tatttatcta cccccccccg ccggtgggag taattgcata cttccatccc     780
tattgcctcg ttttggagga gttggtgact ctcacttcta tcggtaatag gacattaccg     840
tatccgacct tatgactcgg ttccccgatc aacaatcgac tagtaccggc cgcggccacc     900
tacctcctta taacacttct cttaccggca cctccgtcct tggtagtaaa ctcctggcgc     960
tgtatctgtg tgctactgct aggcc                                           985
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<210> 860
<211> 396
<212> DNA
<213> Homo sapiens

<400> 860

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ctgcagaacc gagaggattc ttctgaaggc atcagaaaga agctggtgga agctgaggag      60
ctcgaagaga aacatcggga ggcccaagtc tcagcccagc acctagaagt gcacctgaaa     120
cagaaagagc agcactatga ggaaaagatt aaagtgttgg acaatcagat aaagaaagac     180
ctggctgaca aggagacact ggagaacatg atgcagagac acgaggagga ggcccatgag     240
aagggcaaaa ttctcagcga acagaaggcg atgatcaatg ctatggattc caagatcaga     300
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ctttttaccc aaaggaacat gaaggcccaa tgtatt                               396
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<210> 861
<211> 686
<212> DNA
<213> Homo sapiens

<400> 861

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ccagcaactt tgaaaacatc ctgacgtggg acagcgggcc agagggcacc ccagacacgg     180
tctacagcat cgagtataag acgtacggag agagggactg ggtggcaaag aagggctgtc     240
agcggatcac ccggaagtcc tgcaacctga cggtggagac gggcaacctc acggagctct     300
actatgccag ggtcaccgct gtcagtgcgg gaggccggtc agccaccaag atgactgaca     360
ggttcagctc tctgcagcac actaccctca agccacctga tgtgacctgt atctccaaag     420
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accgcaccta ccaaatggtg agtgtatgtt gcaccctggt ctttctctgc ctaggaagcc     600
tcttccctcc caattagatc tgagttgctt taagaaaaaa aggggacatg ttatgtaaat     660
tagcatttcc cacaacatgt cccttg                                          686
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<210> 862
<211> 383
<212> DNA
<213> Homo sapiens

<400> 862
cagagagttc aagcccacac tccctgggcg tgtctggctg gtgtcacctt ttggagccaa 60
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catggaccaa cagatcacag cagtcatcct caaccgcatg gaatacagac tgcagaaggg 180
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tggactgcct tggtatgtct cagccactgt catctccctg gctcacatgg acagtcttcg 300
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gaggctgaca ggcctggtgg tgt 383

<210> 863
<211> 673
<212> DNA
<213> Homo sapiens

<400> 863
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ggagctggac accaaaggtg agcctggcag gggaggagcg tggggagacc tgtcagcccg 180
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aatgtgttgt ctgttgctag gagacagtct gtaatttacc aaatgtgccg gtccttggcc 360
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ctctgagccc tggtggcttg ggtttagaca gtccccagtg ttgcctgtgt taggggagga 480
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tcaagagaag gtctgaaaac tgaaaagaga gtcctctaag gatccagggt gtccccccac 600
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tcattgataa ggg 673

<210> 864
<211> 435
<212> DNA
<213> Homo sapiens

<220>
<221> misc_feature
<222> (1)...(435)
<223> n = a,t,c or g

<400> 864
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tcatgcgtgc gctgtgtgtt gtgtgtgtat atctgcggag acgcataaag tatgagcgct 120
ttttaggatg ggaattgaga tgtaagattt gggggtgagg gccnccctga cccataggcc 180
tgacatcctc atcctatgga ccctagagtc tggccactcc aggaacctga cctgctctgt 240
gccccgcccc tgtaagcata gaacaccccc catgatctcc tggagtgggg cctccgagac 300

ctccccgggc cccactactg cccgttcctc agtgctcacc cttaccccaa agccccagga 360
nnaccggncc agccctcacc tgtnaggttg accttgcctg gggacagggt gtgacccacg 420
accnatacct ntncg 435

<210> 865
<211> 2161
<212> DNA
<213> Homo sapiens

<400> 865
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agagccgacc gttcaatgtg gctctgaaac tgggccatct ccagagtgga tgctacaaca 120
tgatctaatc ccgggagact tgagggacct ccgagtagaa cctgttacaa ctagtgttgc 180
aacaggggac tattcaattt tgatgaatgt aagctgggta ctccgggcag atgccagcat 240
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ctgtgtgagg tgcaattaca cagaggcctt ccagactcag accagaccct ctggtggtaa 360
atggacattt tcctacatcg gcttccctgt agagctgaac acagtctatt tcattggggc 420
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cctgtgggat ccgaacatca ctgcttgtaa gaagaatgag gagacagtag aagtgaactt 600
cacaaccact cccctgggaa acagatacat ggctcttatc caacacagca ctatcatcgg 660
gttttctcag gtgtttgagc cacaccagaa gaaacaaacg cgagcttcag tggtgattcc 720
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gctggtggcc acatgggtgc tggtggcagg gatctatcta atgtggaggc acgaaaggat 960
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ttgcagaagt gaggtcatcc ttgaaaagtg gcagaaaaag aaaatagcag agatgggtcc 1140
agtgcagtgg cttgccactc aaaagaaggc agcagacaaa gtcgtcttcc ttctttccaa 1200
tgacgtcaac agtgtgtgcg atggtacctg tggcaagagc gagggcagtc ccagtgagaa 1260
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ttagtaatta aaacatttta taccaataaa attttcaaat attgctaact aatgtagcat 1680
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tacagtttta attgaaaact ataaccattt tgataatgca acaataaagc atcttcagcc 1800
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cttaagtgaa aaaccttcta ttacatgcaa aaaatcattg tttttaagat aacaaaagta 1980
gggaataaac aagctgaacc cacttttact ggaccaaatg atctattata tgtgtaacca 2040
cttgtatgat ttggtatttg cataagacct tccctctaca aactagattc atatcttgat 2100
tcttgtacag gtgcctttta acatgaacaa caaaataccc acaaacttgt ctacttttgc 2160
c 2161

<210> 866
<211> 505
<212> DNA
<213> Homo sapiens

<220>

<221> misc_feature
<222> (1)...(505)
<223> n = a,t,c or g

<400> 866
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tggggttgga atattctact ttgttattta tatcatcata tccttcctgg ttgtggtgaa 120
catgtacatt gcagtcatac tggagaattt tagtgttgcc actgaagaaa gtactgaacc 180
tctgagtgag gatgactttg agatgttcta tgaggtttgg gagaagtttg atcccgatgc 240
gacccagttt atagagttct ctaaactctc tgattttgca gctgccctgg atcctcctct 300
tctcatagca aaacccaaca aagtccagct cattgccatg gatctgccca tggttagtgg 360
tgaccggatc cattgtcttg acatcttatt tgcttttaca aagcgtgttt tgggtgagag 420
tggggagatg gattctcttc gttcacagat ggaagaaagg ttcatgtctg caaatccttc 480
caaagtgtcc tatgaaccca tcaca 505

<210> 867
<211> 608
<212> DNA
<213> Homo sapiens

<400> 867
ttcagttttt ggctctggtg caccatgtgc ctgggttaat ttgggtggct caatcccaaa 60
gcagctctga accccaaagc ggctcctctg aattcccagt ttcaagttcc actctgtccc 120
tgctgggcat ctcgagatat gggaaacagg gctgttataa ttgccagaca gctgagttct 180
gtacatacct tgatttgcaa ttttttttgg ctgcttctca ggacaactgg gggagattta 240
gattccttaa aatgcagtta tgaatctatt ggcctcaact ctatttctac ccatgaattc 300
atttgtactt ggcaaagacg acttaatttc tcatttgtta tgtcatttaa acctctcttt 360
agagcctctc ctcactctta cctgttaata atcggaagtc agctacatga aacgttcaat 420
ttgggttcca tctcctctga agaaaaatgc agttaaaaaa aaaataagag gtttggccag 480
ccgcagtggc tcacacctgt aatcccagca ttttgggagg ccgaggcagt cagatcacct 540
gggggcggga gttcgggaac cggcctggcc caacacagga gaaaccccgt cttatactaa 600
acaatata 608

<210> 868
<211> 772
<212> DNA
<213> Homo sapiens

<220>
<221> misc_feature
<222> (1)...(772)
<223> n = a,t,c or g

<400> 868
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ctctggccaa ggtggctggc agacaagatg ctgcccctcc tgggggcagt gctgcttcag 120
aagagagaga agaggggccc tctgtggagg cactggcggc gggaaaccta cccatactat 180
gacctccagg tgaaggtgct gagggccaca aacatccggg gcacagacct gctgtccaaa 240
gccgactgct atgtgcaact gtggctgccc acggcgtccc caagccctgc ccagactagg 300
atagtggcca actgcagtga ccccgagtgg aatgagacct tccactacca gatccatggt 360
gctgtgaaga acgtcctgga gctcaccctc tatgacaagg acatcctggg cagcgaccag 420
ctctctctgc tcctgtttga cctgagaagc ctcaagtgtg gccaacctca caaacacacc 480

ttcccactca accaccagga ttcacaagag ctgcaggtgg aatttgttct ggagaagagc 540
caggagcctg catctgaagt catcaccaac ggggttctgg gggctcaccc ctggctgaga 600
atgaagggta tgattttggg agaggggaga gccccacggc aacagcacgg ccaatcttgg 660
gagggggggg tgggaccctc ccccctctcc ccnngnanaa acaccggagg gaagatagtt 720
gggtttgggg aagaaatggc gaatgggacc ggcgccccac cccgcccccc ct 772

<210> 869
<211> 704
<212> DNA
<213> Homo sapiens

<220>
<221> misc_feature
<222> (1)...(704)
<223> n = a,t,c or g

<400> 869
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tgagccatgg tgtcttctgc cgtcaaaggg cgaccctaac tgcatcctgc tggagtcgag 120
aaaaccaggt agactggaaa ggatgtgtct acagtaactg aaacacatca ctgcgttttg 180
ttacagtcaa tgatagggca gatctgagtt ccagagcacg gctcacagac ctttccttgc 240
atcagtctgt gccgaagtcn nnnnnnnnnc ttttttcttt ttttgcccac attacatcac 300
ttcataattt accacctacg tagcatgact gtatatttgg aatcatttct tcacaagttt 360
tagaccatat taaaggaaca ctggcagaac cctgtttgat ttccctttcg tctgttcccc 420
tacattgccc tcctggcccc cttgaggaac tagatgagcg attagaactg gccagaggtc 480
cttggaggaa caacagcgaa acagaagcat tagtagcatt gtcctcccca gtctaacact 540
tgtcggaccc ctgatgagca gacttccctg tggggtgttc atatccccat gccccgctca 600
gtgggcttca tgtctgagtc atatttgcct gctttccttt gaggtggtgg gcgccaaggt 660
tgtgacaaat gcccggagtc ctggagctcg ctgttacggt tttg 704

<210> 870
<211> 389
<212> DNA
<213> Homo sapiens

<400> 870
tttcgtgagg ctttgttctt ttgttctttg tgatagatct aattgctgct cactctttgg 60
gtctgtactg cgtttatgag ctgtgacact cgccgtgaag gtctgcagct tcactcctga 120
accagcgaga ggaggaaccc accagaagga ggaaaacgcg gaacacatct gaatatcaga 180
aggaacaaac tccagacacg ccgcctttaa gaactgtaac agtcaccgcg agggtccgtg 240
gtttcattct tgaagtaagt gagaccaaga acctgccaat ttcagacaca atggagagcg 300
ccagtcctgc tgcggggcca tacatctatt taatttcctc tcatcttccc cccggttccg 360
agaggaaggt gctttcacct gcactgttc 389

<210> 871
<211> 643
<212> DNA
<213> Homo sapiens

<220>
<221> misc_feature

<222> (1)...(643)
<223> n = a,t,c or g

<400> 871
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aaggagtacg gacgggagtc agaggcagag cgagggtgtg tggagggccg gcggggaccg 120
ccgggagcgc gcggatgtcg gtgttcctgg ggccagggat gccctctgca tctttattag 180
taaatcttct ttcagcttta ctcatcctat ttgtgtttgg agaaacagaa ataagattta 240
ctggacaaac tgaatttgtt gttaatgaaa caagtacaac agttattcgt cttatcattg 300
aaaggatagg agagccagca aatgttactg caattgtatc gctgtatgga gaggacgctg 360
gtgacttttt tgacacatat gctgcagctt ttatacctgc cggagaaaca aacagaacag 420
tgtacatagc agtatgtgat gatgacttac cagagcctga cgaaactttt atttttcact 480
taacattaca gaaaccttca gcaaatgtga agcttggatg gccaaggact gttactgtga 540
caatattatc aaatggacaa atggcatttt gggaatttat tttcatttta aatattggcc 600
ttccccctcc aattccgcca agtggaagnt tgaaagcccc cct 643

<210> 872
<211> 498
<212> DNA
<213> Homo sapiens

<220>
<221> misc_feature
<222> (1)...(498)
<223> n = a,t,c or g

<400> 872
attcccgtgt cgacgatttc gtagcgcctg agagggcggt ggggtggcgg ngttcctgcg 60
cgcggcccgc catggatgtg gaggaggcgt tccaggcggt gggggagatg ggcatctacc 120
agatgtactt gtgcttcctg ctggccgtgc tgctgcagct ctacgtggcc acggaggcca 180
tcctcattgc actggttggg gccacgccat cctaccactg ggacctggca gagctcctgc 240
caaatcagag ccacggtaac cagtcagctg gtgaagacca ggcctttggg gactggctcc 300
tgacagccaa cggcagtgag atccataagc acgtgcattt cagcagcagc ttcacctcta 360
tcgcctcgga gtggttttta attgccaaca gatcctacaa agtcagtgca gcaagctctt 420
ttttcttcag tggtgtattt gttggagtta tctcttttgg tcagctttca gatcgcttcg 480
gaaggaaaaa agtctatc 498

<210> 873
<211> 404
<212> DNA
<213> Homo sapiens

<400> 873
tttcgtctgt gagctgcggc agctgagcag aggcggcggc gcgggacctg cagtcgccag 60
ggattccctc caggtgacga tgctctggtt ctccggcgtc ggggctctgg ctgagcgtta 120
ctgccgccgc tcgcctggga ttacgtgctg cgtcttgctg ctactcaatt gctcgggggt 180
ccccatgtct ctggcttcct ccttcttgac aggttctgtt gcaaaatgtg aaaatgaagg 240
tgaagtcctc cagattccat ttatcacaga caacccttgc ataatgtgtg tctgcttgaa 300
caaggaagtg acatgtaaga gagagaagtg ccccgtgctg tcccgagact gtgccctggc 360
catcaagcag aggggagcct gttgtgaaca gtgcaaaggt tgca 404

<210> 874
<211> 435
<212> DNA
<213> Homo sapiens

<400> 874
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gaattcatcc gtcagtgtgg agtggccctc tgcatcgtgc tgggattctc catcctgtct     60
gcatccatcg gcagctctgt ggtgagggac agggtgattg gagccaaaag gttgcagcac    120
ataagtggcc ttggctacag gatgtactgg ttcacaaact tcctatatga catgctcttt    180
tacttggttt ccgtctgcct gtgtgttgcc gttattgtcg ccttccagtt aacagctttt    240
actttccgca agaacttggc agccacggcc ctcctgctgt cacttttcgg atatgcaact    300
cttccatgga tgtacctgat gtccagaatc ttttccagtt cggacgtggc tttcatttcc    360
tatgtctcac taaacttcat ctttggcctt tgtaccatgc tcataaccat tatgccccgg    420
ttgctagcca tcatc                                                     435
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<210> 875
<211> 703
<212> DNA
<213> Homo sapiens

<400> 875
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cctacttctc cccagtgga tgcagaatgt gctgggccag gtgctggacg cgctggaata     60
cctgcaccat ttggacatca tccacagacc cctttcgtaa gtgctggatg gcccctgaag    120
ccctcaactt ctccttcagc cataaatcag acatctggtc cctgggctgc atcattctgg    180
acatgaccag ctgctccttc atggatggca cagaagccat gcatctgcgg aagtccctcc    240
gccagagccc aggcagcctg aaggccgtcc tgaagacaat ggaggagaag cagatcccgg    300
atgtggaaac cttcaggaat cttctgccct tgatgctcca gatcgacccc tcggatcgaa    360
taacgataaa gtgagctcag ggtcggggtt tattttaacc tgtggattta tctttcaaca    420
tctctccacc ctaatacaag cacagctagt tggctttgta acgcctcaaa gaactccatc    480
acagatgccc tgattatccc tgcacagctg ggctttgccc agttctggct ctcccaaacc    540
gtgctgcggc gagtaatccc gaatgtacgg tggagtgagc agactgaccc ccaggaggca    600
caggaggcgt agcccccagg acccacgaca cttttagggt tccagaaaaa agttttcatt    660
caacataaaa aaaaaaaaat tcctaaagac aaaaaaaaaa aaa                      703
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<210> 876
<211> 429
<212> DNA
<213> Homo sapiens

<400> 876
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tattatgaca gtgcggtgga attcgtggag tgagtctgag gacagcagat gaacagacag     60
aaactgaaag atcccctaat ttgatgagtg agagggtcga gcggaactgg agcacgggcg    120
gctggctgct ggcactgtgc ctggcctggc tgtggaccca cctgaccttg gctgccctgc    180
agcctcccac tgccacagtg cttgtgcagc agggcacctg cgaggtgatt gcggctcacc    240
gctgctgcaa ccggaaccgc atcgaggagc gctcccagac ggtgaaatgc tcctgttttt    300
ctggccaggt ggccggcacc acgcgggcaa agccctcctg cgtggacgac ctgctcttgg    360
ctgcccactg tgctcgtaga gaccctagag ctgcactccg cctcctgctt ccacagcctc    420
catcgtcct                                                            429
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<210> 877
<211> 1140
<212> DNA
<213> Homo sapiens

<400> 877
cgtcactagc agtttctgga gctacttgcc aaggctgagt gtgagctgag cctgccccac 60
caccaagatg atcctgagct tgctgttcag ccttggggc cccctgggct gggggctgct 120
gggggcatgg gcccaggctt ccagtactag cctctctgat ctgcagagct ccaggacacc 180
tggggtctgg aaggcagagg ctgaggacac cggcaaggac cccgttggac gtaactggtg 240
cccctaccca atgtccaagc tggtcacctt actagctctt tgcaaaacag agaaattcct 300
catccactcg cagcagccgt gtccgcaggg agctccagac tgccagaaag tcaaagtcat 360
gtaccgcatg gcccacaagc cagtgtacca ggtcaagcag aaggtgctga cctctttggc 420
ctggaggtgc tgccctggct acacgggccc caactgcgag caccacgatt ccatggcaat 480
ccctgagcct gcagatcctg gtgacagcca ccaggaacct caggatggac cagtcagctt 540
caaacctggc caccttgctg cagtgatcaa tgaggttgag gtgcaacagg aacagcagga 600
acatctgctg ggagatctcc agaatgatgt gcaccgggtg gcagacagcc tgccaggcct 660
gtggaaagcc ctgcctggta acctcacagc tgcagtgatg gaagcaaatc aaacaggcca 720
cgagttccct gatagatcct tggagcaggt gctgctaccc cacgtggaca ccttcctaca 780
agtgcatttc agccccatct ggaggagctt taaccaaagc ctgcacagcc ttacccaggc 840
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cagtgccgtg gccagggctg acttccagga gcttggtgcc aaatttgagg ccaaggtcca 960
ggagaacact cagagagtgg gtcagctgcg acaggacgtg gaggaccgcc tgcacgccca 1020
gcactttacc ctgcaccgct cgatctcaga gctccaagcc gatgtggaca ccaaattgaa 1080
gaggctgcac aaggctcagg aggccccagg gaccaatggc agtctggtgt tggaacgcct 1140

<210> 878
<211> 1139
<212> DNA
<213> Homo sapiens

<400> 878
tgccactgtg aaggagatga tgagagcccc ctgatcaccc cctgccactg cacaggaagc 60
ctccacttcg tgcaccaggc ctgcctgcag cagtggatca agagctccga cacgcgctgc 120
tgcgagctct gcaagtatga gttcatcatg gagaccaagc tgaagccact gagaaaatgg 180
gagaagttgc agatgacgtc cagcgagcgc aggaagatca tgtgctcagt gacattccac 240
gtcattgcca tcacatgtgt ggtctggtcc ttgtatgtgc tcattgaccg tactgctgag 300
gagatcaagc aggggcaggc aacaggaatc ctagaatggc ccttttggac taaattggtg 360
gttgtggcca tcggcttcac cggaggactt ctttttatgt atgttcagtg taaagtgtat 420
gtgcaattgt ggaagagact caaggcctat aatagagtga tctatgttca aaactgtcca 480
gaaacaagca aaaagaatat ttttgaaaaa tctccactaa cagagcccaa ctttgaaaat 540
aaacatggat atggaatctg tcattccgac acaaactctt cttgttgcac agagcctgaa 600
gacactggag cagaaatcat tcacgtctga ttgtgtgcgg gttgtcattt tcctggacat 660
ccatgaagag ctgaaggaaa ttgtttactg ccaattgtat acctttctta tgtcctttaa 720
tagcatagac tggacaggtg actatttata gtggcttctc tttttctaaa ccctccttag 780
tctcctagaa aaccttcctg tgggccaggc atgcctgggt cctgcctctg cctggcagct 840
ctgtgggaaa gtggaagacc ccatgatgac atcatgggga gccagcagag ttcctgccca 900
tggtcttgag ctgaatgaga gaataaaatg ccaatcccaa gggaagagga ggagcagggg 960
tgcccaggcc ctgataccca gccgcctcca gcttgcagtg gtccccagcc tggagcagag 1020
cattggggag tgtctaagcc atgacgagaa gattccctct gcatcacggc gaaccccccag 1080
gagatggtat ttgaaaacag acccccaaac acagactcct gcctgccctc ttgccgatg 1139

<210> 879

<211> 478
<212> DNA
<213> Homo sapiens

<220>
<221> misc_feature
<222> (1)...(478)
<223> n = a,t,c or g

<400> 879
ggtcacgcaa gcggcnncnn nttttgagac ctttgatagc gtgtaggaan ncccaggcca 60
gtgaatgtca gttcgtcggg cactgactcc gtctgctctt ggccttgtgt tcattttaca 120
aatatttgcc cacggcctcc caggcccagg cccatgccac ctgggccccg gcatctgttt 180
gaggatctgc caatgtgctc ttaactgagg acgaaggaag aacacctttc tatgagtctt 240
gcaaagatta cctccttcag gccacaaata tttgagtgca cactacgtgc caggcactgt 300
gcagggctgc aggcatagag acagaatgta atctatctgg gccttggacc ccatagggag 360
aggggaccac tcaggtccat acttcctttg gacttggggc tttggccttg ggagggggcgg 420
aggtggcgtg gcaagatgaa aaagacatcc tgcccccatc cacttgggca gagcttct 478

<210> 880
<211> 546
<212> DNA
<213> Homo sapiens

<400> 880
atgctgggta tccgtgatgt gagagggttt agcacgggaa cactgcagac gcctgcctgg 60
gagctcaggt gctgcggtcc tcccttctgc ctgaaggagg catatggcca ggggctccgc 120
ctgacactca cgaggcagta tatgcggatg atgggagtgc atccagtgat ccatttcctg 180
gcctggttcc tggagaacat ggctgtgttg accataagca gtgctactct ggccatcgtt 240
ctgaaaacaa gtggcatctt tgcacacagc aataccttta ttgtttttcct ctttctcttg 300
gattttggga tgtcagtcgt catgctgagc tacctcttga gtgcattttt cagccaagct 360
aatacagcgg ccctttgtac cagcctggtg tacatgatca gctttctgcc ctacatagtt 420
ctattggttc tacataacca attaagtttt gttaatcaga catttctgtg ccttctttcg 480
acaaccgcct ttggacaagg ggtatttttt attacattcc tggaaggaca agagacaggg 540
attcac 546

<210> 881
<211> 918
<212> DNA
<213> Homo sapiens

<400> 881
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tttgctctga ttgaacggaa tgttccaccg tgtttcatct ttattcatta tcctttgttc 120
tttaaaatct gatatattgg cataaaagta attgtacata tatatatgaa tgtgatttat 180
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tgcacaaatc tgtgtgcagt gctttttgcc cgttgcctag acgatcactt ggtttctctg 300
aggatgtctg gttctcgtaa agagtttgat gtgaaacaga ttttgaaaat cagatggagg 360
tggtttggtc atcaagcatc atctcctaat tctacagttg acagccagca gggagaattt 420
tggaaccgag gacagactgg agcaaacggt gggagaaagt ttttagatcc atgtagccta 480
caattgcctt tggcttcaat tggttaccga aggtccagcc aactggattt tcagaattca 540
ccttcttggc caatggcatc cacctctgaa gtccctgcat ttgagtttac agcagaagat 600

tgtggcggtg cacattggct ggatagacca gaagtggatg atggcactag tgaagaagaa 660
aatgaatctg attccagttc atgcaggact tccaatagta gtcagacatt atcatcctgt 720
catactatgg agccatgtac atcagatgaa tttttccaag cccttaatca tgccgagcaa 780
acatttaaaa aaatggaaaa ctatttgaga cataaacagt tgtgtgatgt aattttagtc 840
gctggtgatc gcagaattcc agctcacaga ttggtgctct cctctgtctc agactatttt 900
gctggcatgt ttactaat 918

<210> 882
<211> 604
<212> DNA
<213> Homo sapiens

<220>
<221> misc_feature
<222> (1)...(604)
<223> n = a,t,c or g

<400> 882
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atgtgatgga cacgctgggg ctttttttact tcatagcagg aattgtattt cggctccact 120
cttctaataa aagctctttg tattctggac gagtcatttt ctgtctggac tacattattt 180
tcactctaag attgatccac atttttactg taagcagaaa cttaggaccc aagattataa 240
tgctgcagag gatgctgatc gatgtgttct tcttcctgtt cctctttgcg gngtggatgg 300
tggcctttgg cgtggccagg caagggatcc ttaggcagaa tgagcagcgc tggaggtgga 360
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tggatggtac cacgtatgac tttgcccact gcaccttcac tgggaatgag tccaagccac 480
tgtgtgtgga gctggatgag cacaacctgc cccggttccc cgagtggatc accatccccc 540
tggtgtgcat ctacatgtta tccaccaaca tcctgctggt caacctgctg gtcgccatgt 600
ttgg 604

<210> 883
<211> 1206
<212> DNA
<213> Homo sapiens

<400> 883
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cctacagggc ctgctggaga agaccaatgg gtgcatggga tgaccggcag cttccctcaa 120
gtggcttccc agagactact aggagaactt ggtcctatcg ctgcccccac ctggaagctg 180
gacttaagga tcccccaaag aacggggcaa ttagaaacct cccacccagc gaagggataa 240
gcttctcaac tcagtcccac cactcttcat cgcaaccctc tgagtctgca gcagaaacaa 300
acatctccaa gttacagagg aggggatgga atccccaagg ggccgagcgg tagccctttt 360
aacttataag cctgttgatt agcctatacg agttatttgc acgtcaagaa aggaagtagc 420
ctgctccttc ctgcagcgtc ctgctggtgt gacagcacgt ccccaagctc agtgctaacc 480
tccttattaa acatcccctg ctgtgactca gggaacccac atgggtactc taaaacagtc 540
attcagggac cccacggggt catgtgggag ggagacagat cccagaaaga gcacaagtga 600
gtcattacca aaaactccaa ggcccgcaca ccggacgcac atacccagct aggggcagac 660
tcaaagatcc cagcccttat cttctcccca tatcagagct cggaagccag aaatcttcct 720
aaggcaggtg aaagcaagcc gagcccccact gctgaaggac aaagccacag gaagcctgat 780
gacatctttc ctctgaggct tccaaacgat caccccaaat tgcttgctga tactgggaag 840
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atttaaggag aaagtcaaag tagccaaaag gcaaaccaga tggtggtgga catgtgggtg 960
acagagcatc ctgcatttgt tgcctcgggg tgcagcccca aagataaagc cagcagtgtg 1020

caaatgacaa atgctacccc acctccgcca ggcagccaga gccagggccg aaggacgcgg 1080
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ctcgat 1206

<210> 884
<211> 420
<212> DNA
<213> Homo sapiens

<220>
<221> misc_feature
<222> (1)...(420)
<223> n = a,t,c or g

<400> 884
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cggtgcagca gtattcgccc aacgggcgta tcggaaacca ctgatctcgt tccttgtcgg 180
cttctcgatg ctggcggccg gcgtaaccag tgcggcggga ctcgccctcg ccttctcggg 240
cgactatctc aaagccttca tcgacgtccc aaccgttcca gcggcgctcg tcttcctgct 300
cctggtggga cttctcaatg ccagaggcat caaggagtcc atgcgcgcca ncgtcgtcat 360
gacagtcgtg gaagtcaccg ggctcgtcct cgttgtcgtc ctcgcgctcg tgccaggcag 420

<210> 885
<211> 1696
<212> DNA
<213> Homo sapiens

<400> 885
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ctgaactccc ccatccatca gagggtaatg atgagaaggc tgaagaggcg ggtgaaggtc 420
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ctgcaaaaaa aaaaaa 1696

<210> 886
<211> 1410
<212> DNA
<213> Homo sapiens

<400> 886
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gcgtgtggag tatctgcaga catgactgcg tggaggagat tccagtcgct gctcctgctt 240
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ggcttagtcc agggtcccaa tggcactatt gagagcccag ggtttcctca cgggtatccg 360
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ttccatacct ttgctcttga agaagatttt gatattttat cagtttacga tggacagcct 480
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aaagcattat atgaaggtag gagattggtt gtgttttgca catgcattca ctgtccaaat 660
gatctaatac atgctacact ggattaataa tgacaaacta ggctgctatg tcgcaggtcg 720
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atcgccaaac tttccagccc aaaacgtgtt cacaattttc ttcttatcac tagaactttt 840
cttcttgggg ttttgttttg gttaatttgt agcgaataag ttttgagaaa tttgactata 900
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ttaatacttt aaaggccgat gtgtgtctac ttttgttttg catggagatt ttaaattgcc 1320
ttttacacgt aatacaagag ctactgtctg taacagaaac tctggagtct gtaaatttaa 1380
aaagcaatct atcgttaggg gtgctgtatt 1410

<210> 887
<211> 413
<212> DNA
<213> Homo sapiens

<400> 887
tgactcccag aacaaccagt atattttgac caaacccaga gattcaacca tcccacgtgc 60
agatcaccac tttataaagg acattgttac cataggaatg ctgtctttgc cttgtggctg 120
gctatgtaca gccataggat tgcctacaat gtttggttat attatttgtg gtgtacttct 180
gggaccttca ggactaaata gtattaaggt aagaacaaaa ttggattgtt ttggtatctg 240
tttaacagaa tataaaaaga gaattcatga agactaaaaa gtattgaatg tgattaatgc 300
agataccagc ttcgtataaa ccatttcaaa gatgtccttt caggtgtcac gggaagtctc 360
tgaaccctca ggaagtcgct gtgcctgtta gtgaaggggc ggtgttactg gaa 413

<210> 888
<211> 887
<212> DNA
<213> Homo sapiens

<220>
<221> misc_feature
<222> (1)...(887)
<223> n = a,t,c or g

<400> 888
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ataggtcaag taagtaaata gagatttaaa aaattatgaa cacaaaggaa gtaacagcct    180
tcctgtcttg ctgtagtaac tgaccatatg cgtttatatc atgctaattg tgcaatttat    240
ttttgagttg gtctcaagta ttttggtttc gaatgtgaaa gatatgttag atttgaaag     300
tggtttttgt agtaaaattc tcagttattt tttttcttcg ccaagataca gattaccttt    360
cctttaagct gatcctaagg aagttatttt ttgtatacct tcagagaggg gataacatcc    420
caaagatatt agtgttcaca gaggatggat atttcctacg agcctggaat tatacagttg    480
acacacctca tggtatattt gcagccagta ctctatatga acaatccgtc tggatcacgg    540
atgtaggaag tggattcttt ggtcatactg ttaaaaaata cagttctttt ggtgatcttg    600
ttcaagtctt gggtactcca ggcaaaaaag gcactagttt gaatcctttg cagtttgata    660
acccagcaga attatatgta gaggacacag gagatattta cattgtggat ggagatggag    720
gattgaataa cagattgatc aaactgtccc aagatttcat gatcctttgg ctgcatggag    780
aaaatgggac agggcctgct aagttcaaca tacctcacag tgttacactt gattcagctg    840
gtcgggtaca aatacagcgt cattgtgtct gggaaaaaaa aaaaaaa                 887
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<210> 889
<211> 1871
<212> DNA
<213> Homo sapiens

<400> 889
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gcagccttca gcccagtgcc ccctactgag gccaaagcgg caggacccag gccttctggc    120
ctccctgacc tgctcacctc cacgcggctg gccacacacg tctgccaacc ctttcctgtg    180
ccgggggggt ttctccccaa gccctggggc cagctcctcc aagacgctct gcccaccagt    240
ctcaccggac ttggtgaaca ggggcagctc aggattaggg actccctgga cccacccgaa    300
gttctaaggc gggggcccg tgtccccaca gagcctggcc tggagccctg gaaggaggcc    360
ctggtgcggc cccaggcag ctacagcagc agcagcaaca gtggagactg ggatggac      420
ctggccagtg accagtcctc tccgtccacc ccgtcacccc cactgccccc cgaggcagcc    480
cactttctgt ttggggagcc caccctgaga aaaaggaaga gcccggccca ggtcatgttc    540
cagtgtctgt ggaagagctg cgggaaggtg ctgagcacgg cgtcggcgat gcagagacac    600
atccgcctgg tgcacctggg gaggcaggca gagcctgatc agagtgatgg tgaggaggac    660
ttctactaca cagagctgga tgttggtgtg gacacgctga ccgacgggct gtccagcctg    720
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gccttgtcct ccaggatcgg agtcaccctg aggaagcccc gcggcgacgc gaagaagtgc   1020
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gcctgccagc ggttcctgga ctaagtccgg ctcgttcaag aacataagct accaccttct   1140
ccctccccac cccctccagg cccggggctg aaacagcccg aggacagccc caggggctgg   1200
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cctccccccc gccaggtcgg ggaggggtcc caccactcaa agtgcctcta aagaaaccag 1320
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gccaccctcc cgcctgtcgg cccgtagatt tatcaagggt gttatgggcc cagctttggg 1440
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aatttattga aagtgatcgc tttgcaagga tgtctaagct aatcccgtca cagaaaggaa 1860
acgcacaggc g 1871

<210> 890
<211> 379
<212> DNA
<213> Homo sapiens

<400> 890
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gacttgtctt cggtagggac agtcaagtca ggcaaaaccg tgaacttggc tacagcaggc 180
acaatcaagc cgggcacagc catgaatctg actacagttg ggacaaccaa gccagggatg 240
gtcatggatt tgatagcctc agaaccagac aagctgggca aagccatggc tacaagaagc 300
acagccaaac cagatatgac cacagagggt atagccatgg attcagcaac atcagaccca 360
gtcaagccgg acatgtatt 379

<210> 891
<211> 397
<212> DNA
<213> Homo sapiens

<400> 891
tgctgcacaa catgcgtgtg tacggcacgt gcacgctcgt gctcatggcc ctggtggtct 60
tcgtgggcgt caagtatgtc aacaagctgg cgctggtctt cctggcctgc gtcgtgctgt 120
ccatcctggc catctatgcc ggcgtcatca agtctgcctt cgaccccccg gacatcccgg 180
tctgcctcct ggggaaccgc acgctgtcac ggcgcagctt cgatgcctgc gtcaaggcct 240
acggcatcca caacaactca gccacctccg cgctctgggg cctcttctgc aacggctccc 300
agcccagcgc cgcctgtgac gagtacttca tccagaacaa cgtcaccgaa attcagggca 360
tcccgggcgc ggccagtggt gtcttcctgg agaaccg 397

<210> 892
<211> 398
<212> DNA
<213> Homo sapiens

<400> 892
cctgtccgag tccctgctcc tggtcattgc tgacctgctc ttctgccggg acttcacggt 60
tcagagcccc cggaggagca ctgtggactc ggcagaggac gtccactccc tggacagctg 120
tgaatacatc tgggaggttg gtgtgggctt cgctcactcc ccccagccta actacatcca 180
cgatatgaac cggatggagc tgctgaaact gctgctgaca tgcttctccg aggccatgta 240

cctgcccccca gctccggaaa gtggcagcac caacccatgg gttcagttct tttgttccac 300
ggagaacaga catgccctgc ccctcttcac ctccctcctc aacaccgtgt gtgcctatga 360
ccctgtggaa tacgggatcc cctacaacca cctgtatt 398

<210> 893
<211> 397
<212> DNA
<213> Homo sapiens

<400> 893
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tggtccgggg gctgttgctg cgaggggcag ttgatggcat cctacacatg ttcactccca 120
agctggacaa gatgctggac ccccaggtgt ggcgggaggc agctacccag gtcttctctg 180
ccttgggcct gggctttggt ggtgtcattg ccttctccag ctacaataag caggacaaca 240
actgccactt cgatgccgcc ctggtgtcct tcatcaactt cttcacgtca gtgttggcca 300
ccctcgtggt gtttgctgtg ctgggcttca aggccaaçat catgaatgag aagtgtgtgg 360
tcgagaatgc tgagaaaatc ctagggtacc gtgtatt 397

<210> 894
<211> 380
<212> DNA
<213> Homo sapiens

<400> 894
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atggaaccaa gaagttcatg caggagctga cggagatgct gggcttccgc ccctaccgct 120
tctatttcta catgtggaag ttcgtgtctc ctctatgcat ggctgtgctc accacagcca 180
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ccgagcgcta cctgtatttc cccaactggg ccatggcacc cctgatcacc ctcatcgtcg 300
tggcgacgct gcccatccct gtggtgttcg tcctgcggca cttccaccta atctgtgatg 360
gctccaacac cccatgtatt 380

<210> 895
<211> 389
<212> DNA
<213> Homo sapiens

<220>
<221> misc_feature
<222> (1)...(389)
<223> n = a,t,c or g

<400> 895
ncatgaagat gtttgtggct catgggttct atgctgccaa attcgtagtg gccattgggt 60
cggttgcagg actgacagtc agcttgctgg ggtccctctt cccgatgccg agggtcattt 120
atgccatggc tggtgacggg ctcctttca ggttcctggc tcacgtcagc tcctacacag 180
agacaccagt ggtggcctgc atcgtgtcgg ggttcctggc agcgctcctc gcactgttgg 240
tcagcttgag agacctgata gagatgatgt ctatcggcac gctcctggcc tacaccttgg 300
tctctgtctg tgtcttgctc cttcgacacc accctgagag tgacattgat ggttttgtca 360
agttcttgtc tgaggagcac acgtgtagt 389

<210> 896
<211> 415
<212> DNA
<213> Homo sapiens

<400> 896
cagcagccca cctggagtgc atttttaggt ttgaattgag agaacttgac cctgaggcac 60
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caaatgggcc aaagatgggc ctcctgatga tgattctagg ccaaatattc ctgaatggca 180
accaagccaa ggaggctgag atttgggaaa tgctctggag gatgggggtg cagcgggaaa 240
ggaggctttc catttttggg aacccaaaga gacttctgtc tgtggagttt gtatggcagc 300
gttacttaga ctacaggcca gtaactgact gtaaaccagt ggagtatgag tttttctggg 360
gcccaagatc ccacctagaa accaccaaga tgaaaattct gaagttcatg gcgaa 415

<210> 897
<211> 428
<212> DNA
<213> Homo sapiens

<400> 897
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agaaagagtt tgttgctcag cccaactgcc aacagttgct tgccaccctg tggtatgatg 180
gcttccctgg atggcggcgg aaacactggg tagtcaagct tctaacctgc atgaccattg 240
ggttcctgtt tcccatgctg tctatagcct acctgatctc acccaggagc aaccttgggc 300
tgttcatcaa gaaacccttt atcaagttta tctgccacac agcatcctat ttgaccttcc 360
tctctatgct tctcctggct tctcagcaca ttgtcaggac agaccttcat gtacaggggc 420
cctgtatt 428

<210> 898
<211> 444
<212> DNA
<213> Homo sapiens

<220>
<221> misc_feature
<222> (1)...(444)
<223> n = a,t,c or g

<400> 898
ncttacaatg cacaatatct gcttcacatc cttgcccatc ctggcctata gtctactgga 60
acagcacatc aacattgaca ctctgacctc agatccccga ttgtatatga aaatttctgg 120
caatgccatg ctacagttgg gccccttctt atattggaca tttctggctg cctttgaagg 180
gacagtgttc ttctttggga cttactttct ttttcagact gcatccctag aagaaaatgg 240
aaaggtatac ggaaactgga cttttggaac cattgttttt acagtcttag tattcactgt 300
aaccctgaag cttgccttgg atacccgatt ctggacgtgg ataaatcact ttgtgatttg 360
gggttcttta gccttctatg tatttttctc attcttctgg ggaggaatta tttggccttt 420
tctcaagcaa cagagaatgg cgaa 444

<210> 899
<211> 436
<212> DNA
<213> Homo sapiens

<400> 899
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tctccagcaa caggtcttaa acagtgggtg gaagctgtac agggataccc aggatgggga 120
agcctttcaa ggtgaacaga atgatttcaa ctccagccaa ggtgggaaag acttttgcca 180
ccaacatggg ctgtttgagc accaaaaaac ccataatggg gagaggcctt atgagttcag 240
tgaatgtggg gaattgttta ggtacaactc caaccttatt aaatatcagc aaaatcatgc 300
tggagaaagg ccttatgagg gcactgaata tggaaagacc tttattagaa agtccaacct 360
agttcagcac cagaaaattc acagtgaagg ctttctttca aaaaggtctg accccattga 420
acatcaggag tgtatt 436

<210> 900
<211> 466
<212> DNA
<213> Homo sapiens

<220>
<221> misc_feature
<222> (1)...(466)
<223> n = a,t,c or g

<400> 900
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ctgttgggcc tccgagagga ctgggatgac cgctggatca acgatgtgga agacagctac 180
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tcggtcttcc agccggggat gaagaacaag atcttgatat ttggcctctt tgaagagaca 360
gccctggctg ctttcctttc ctactgccct ggaatgggtg ttgctcttaa gatgtatccc 420
ctcaaaccta cctggagggt ctgtgccttc ccctactctc ttctca 466

<210> 901
<211> 412
<212> DNA
<213> Homo sapiens

<400> 901
caagatctgg atcggcccca atgatgggat cacccagttt gataacatcc tttttgctgt 60
gctgactgtc ttccagtgcg tccccatgga agggtggttc gctgtgctgt caactgccaa 120
tgatgtctta ggagccccct ggaattggct gtacttcatc cccctcctca tcattggagc 180
cttctttgtt cccaccctag tcctgggagt gctttccggg gattttgcca aagagagaga 240
gagagtggag acccgaaggg ctttcatgaa gctgcggcgc cagcagcaga ttgagcgtga 300
gctgaatggc taccgtgtct ggatagccaa agcagaggaa gtcatgctcg ctgaagaaaa 360
tttgtatccc agtcacgcac ggccagtgaa tccgtaatca tggtcataga cc 412

<210> 902
<211> 1334
<212> DNA
<213> Homo sapiens

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<211> 1219
<212> DNA
<213> Homo sapiens

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<211> 814
<212> DNA
<213> Homo sapiens

<220>
<221> misc_feature
<222> (1)...(814)
<223> n = a,t,c or g

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<211> 687
<212> DNA
<213> Homo sapiens

<220>
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<223> n = a,t,c or g

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<210> 914
<211> 620
<212> DNA
<213> Homo sapiens

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<210> 915
<211> 788
<212> DNA
<213> Homo sapiens

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<210> 916
<211> 758
<212> DNA
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<210> 917
<211> 2709
<212> DNA

<213> Homo sapiens

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<210> 918
<211> 1327
<212> DNA
<213> Homo sapiens

<220>

<221> misc_feature
<222> (1)...(1327)
<223> n = a,t,c or g

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<210> 919
<211> 1463
<212> DNA
<213> Homo sapiens

<400> 919
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<212> DNA
<213> Homo sapiens

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tcccgccggg gccgacccga cctatagaac tcccccgccc c 761

<210> 921
<211> 1225
<212> DNA
<213> Homo sapiens

<400> 921
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<212> DNA
<213> Homo sapiens

<220>
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<223> n = a,t,c or g

<400> 922
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tgcggagaac actggattta aataaaggtg ttatgggtat aattaactat aatttatttt     180
atgaataaat aagaaaaaag tccctggcta taaaggatag tggaggttta tttcccaata     240
ttcttccatt tggactgaac ccctgccaac taagcagaca tcccactatt acaccaaaaa     300
gactgctctc cctgcgagca attacaatac atgacaccac taggcccaat ggctcagagg     360
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<210> 923
<211> 1071
<212> DNA
<213> Homo sapiens

<400> 923
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ttggtaattt ttagtggatt cggtgtttca ccatgttggc cgggcttgtc ttgaactcct     180
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<210> 924
<211> 1758
<212> DNA
<213> Homo sapiens

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<210> 925
<211> 854
<212> DNA
<213> Homo sapiens

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<210> 926
<211> 2422
<212> DNA
<213> Homo sapiens

<400> 926
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<210> 927
<211> 415
<212> DNA
<213> Homo sapiens

<220>
<221> misc_feature
<222> (1)...(415)
<223> n = a,t,c or g

<400> 927
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ctgaacgact cactttagct tttcttatag gacaggtcta ctagtgacaa actctgtcca    180
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<211> 1503
<212> DNA
<213> Homo sapiens

<220>
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<222> (1)...(1503)
<223> n = a,t,c or g

<400> 928
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<210> 929
<211> 834
<212> DNA
<213> Homo sapiens

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<210> 930
<211> 1434
<212> DNA
<213> Homo sapiens

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<211> 2361
<212> DNA
<213> Homo sapiens

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<212> DNA
<213> Homo sapiens

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<212> DNA
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<212> DNA
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<210> 941
<211> 1510
<212> DNA
<213> Homo sapiens

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<210> 942
<211> 2226
<212> DNA
<213> Homo sapiens

<400> 942
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<210> 943
<211> 1026
<212> DNA
<213> Homo sapiens

<400> 943
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aacaactgaa acatatgtca acagtaaata aactgcagta tattcataca atgatttact 240
ctatggcaat gacaataaac aaactatatg caaaacatgg atgaccctta caagccttag 300
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<212> DNA
<213> Homo sapiens

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ttatgattac atccaatatt ttgaaacaca tttgttgtca tctaaaagtt cgacagacta 240
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acactctcgt gggttttggt agtagagctg gaagtcaaag ccaagtcagt ctttttattg 480
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<210> 945
<211> 2127
<212> DNA
<213> Homo sapiens

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<210> 946
<211> 1759
<212> DNA
<213> Homo sapiens

<400> 946
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<210> 947
<211> 1033
<212> DNA
<213> Homo sapiens

<220>
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<222> (1)...(1033)
<223> n = a,t,c or g

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<210> 948
<211> 401
<212> DNA
<213> Homo sapiens

<400> 948
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gctggtctgc tcgggccacg acggattcct gggctttcat gggtggggca gcccgtgtcc 360
cggccccccg ttccggtgca acactctgga tgcggagggc g 401

<210> 949
<211> 432
<212> DNA
<213> Homo sapiens

<400> 949
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ggtcgcgatg aa 432

<210> 950
<211> 450
<212> DNA
<213> Homo sapiens

<400> 950
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ggacaagtga aattctctga gagccattgg tcagtacaat gaatatgaaa ttcatgcctg 180
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<210> 951
<211> 1321
<212> DNA
<213> Homo sapiens

<220>
<221> misc_feature
<222> (1)...(1321)
<223> n = a,t,c or g

<400> 951
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<210> 952
<211> 1729
<212> DNA
<213> Homo sapiens

<400> 952
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taaaaatgta aaggggagag tgggtacata tctgaacatt aaactttagg cactttctgg    240
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gccgacccgc cgctgccgcc gctcccctcg cgctccatcc cgtcgccatt caccacagag   1680
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<212> DNA
<213> Homo sapiens

<400> 953

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<212> DNA
<213> Homo sapiens

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<223> n = a,t,c or g

<400> 954
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<211> 1172
<212> DNA
<213> Homo sapiens

<400> 955
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<210> 956
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<212> DNA
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<400> 956
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<211> 2874
<212> DNA
<213> Homo sapiens

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<212> DNA
<213> Homo sapiens

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<211> 476
<212> DNA
<213> Homo sapiens

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<400> 959
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<211> 679
<212> DNA
<213> Homo sapiens

<400> 961
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ccctcccctt gtcgccccc 679

<210> 962
<211> 782
<212> DNA
<213> Homo sapiens

<220>
<221> misc_feature
<222> (1)...(782)
<223> n = a,t,c or g

<400> 962
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gtcctgattt gcagggtgtc ctagtggtga tgctcatctt cactgtgctg gagctcttat 180
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<210> 963
<211> 1734
<212> DNA
<213> Homo sapiens

<220>
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<222> (1)...(1734)
<223> n = a,t,c or g

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<210> 964
<211> 1098
<212> DNA
<213> Homo sapiens

<400> 964
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aggcatgggc catcatgcct ggctaatttt tatattttta atggagacat ggtttcacta 180
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caggtcctcg acccggga 1098

<210> 965
<211> 422
<212> DNA
<213> Homo sapiens

<400> 965
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tcaacgactg ccgagacatc attcagcgca tgcaccttcg tcagtacgag ctgctctaaa 300
aagggaaccc ccagatttag ttaaagcctt aggcacaatt agttaaaagt gaaacgtaat 360
tgtacaagca gttaatcacc caccataggg catgagtaac aaagcaacct ttcccttacc 420
cg 422

<210> 966
<211> 617

<212> DNA
<213> Homo sapiens

<400> 966
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gcagtctctc agaaagagac acacccaccc cacaaaaaaa tccttggaat ttgtatgtgc 180
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<210> 967
<211> 1446
<212> DNA
<213> Homo sapiens

<400> 967
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gagtacctga ttttgcagcc attaaaaatg tgttcattaa ctatgtgggg aattattgga 1440
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<210> 968
<211> 1495
<212> DNA
<213> Homo sapiens

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taaatttccg ggccggggtt taaaaaacgc ggggagggaa aaaccccggg cttcc 1495

<210> 969
<211> 999
<212> DNA
<213> Homo sapiens

<400> 969
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tcatcatctc cggatccctc tcagtggcag ccgagaagaa ccacaccagt tgcctggtga 180
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<210> 970
<211> 865

<212> DNA
<213> Homo sapiens

<400> 970
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<210> 971
<211> 630
<212> DNA
<213> Homo sapiens

<400> 971
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taacatttat ttgtattgcc cacctgtatt 630

<210> 972
<211> 426
<212> DNA
<213> Homo sapiens

<400> 972
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aaaagc 426

<210> 973
<211> 542
<212> DNA
<213> Homo sapiens

<400> 973
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ga 542

<210> 974
<211> 2870
<212> DNA
<213> Homo sapiens

<400> 974
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gcctgtaatt cagtgtttac agcattagat cactgtcatg aagccataga aataacaagc 180
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aataagcaga ttcacaagat tcatcgtgat tcaggagata attctcagac agagcctcat 540
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<212> DNA
<213> Homo sapiens

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<210> 984
<211> 1813
<212> DNA
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<212> DNA
<213> Homo sapiens

<400> 985

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<210> 986
<211> 876
<212> DNA
<213> Homo sapiens

<400> 986

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<211> 1884
<212> DNA
<213> Homo sapiens

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<210> 988
<211> 935
<212> DNA
<213> Homo sapiens

<400> 988

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<210> 989
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<212> DNA
<213> Homo sapiens

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<212> DNA
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<212> DNA
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<211> 447
<212> DNA
<213> Homo sapiens

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<210> 993
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<212> DNA
<213> Homo. sapiens

<400> 993

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<212> DNA
<213> Homo sapiens

<400> 994

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<212> DNA
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<212> DNA
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<212> DNA
<213> Homo sapiens

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aggcccaggc gatcctcccc cctcctgcct cc 752

<210> 1008
<211> 1145
<212> DNA
<213> Homo sapiens

<400> 1008
caatgatatg ctcttctaca ctcttaaaaa ttatagacaa ccccaaataa cttttattta 60
gtggttttaa caatatttac catgtctgaa atatgataaa cattaaaatt agtattttgg 120
aaaaatgcca tattagaaac tgatgattta aaagtaacaa caatgaatcc attacatgtg 180
aacatactgt tttttgtttt gtttgtttgt ttgttttgag acggagtttc actcttttgc 240
ccaggctgga gtgcagtggt gcgattgcag ctcactgtag tcttcgcctc ccaggctcaa 300
gtgattctca tgcctcagcc tcctgagtag ctgggattac aggtgctcac caccacaccc 360
ggctaatttt tgtagagatg gggtttcacc gtattggcca ggctggtctt gaactccaga 420
cttcaagtga tccacccacc ttggcctccc aaagtgctgg gattacgggc atgagccact 480
gcaccaggcc aacatacttt ttataaaaac agctgtcttc tctaaaacaa caaaaaaatg 540
tagataatag tagtatcatt ttatagtttt gcaactctct ttaatgtttg gcttaataaa 600
agatagttgg attctcgtat ctgtttttgt attcagtctg ttgtggatgg tgatttgatt 660
gaagtaaatg aaggaaatcc agctacatac agatttggag ttggaaaaaa tagtatttta 720
ataacctttt tagatcatgg tggatactct tcttttgttt ggcctcaaaa ttagaacaaa 780
ggcagtttct gaaaataatt gtatgtggtg aaaaattaat gaatcttata tggaccatac 840
ttttaattta gaatattggt ctaaaaaaaa aaaagggggc cctttaaaaa caaatttagt 900
acgggcgtgg atgttaactt ttttggggcc agattgttcg ggcgggtgta caggggaagg 960
ggaaaacggg tggggctagg acgtgttgaa caaatgacgt gctcgtgctg gcgaccgacc 1020
tcttgtacga gaggtaatgc gattgggaac gagtgatggg tgcgtcgatt ggtcgaggcg 1080
tgcgatgcat gcaatggggc gcttaggcgt tgggtaggat gggtgggacg gatcgaacgt 1140
tctcg 1145

<210> 1009
<211> 737
<212> DNA
<213> Homo sapiens

<400> 1009
actgtatatc taaattgcag tgtcacccct cccatgctcc acatttcttc agcctttcac 60
tgctatgctt ttcttccact ttttgctctg acacataatt tcatttttctt attttattta 120
ttatctctct cccccaaact agaatgtaaa ttccaggaag gcagagattt ctatctattt 180
ttttttgtct tccccatatt ctggcatgtc tggcatagaa aaggcattta gtaaacattt 240
gttaaatgaa ttgactatct tttctctgca aacttgttcc tcaaattctg ccaaacctaa 300
attgaaacaa gcaggtattg tattttggta caagtcctgg ggctgtggat taaatccaag 360
agcattgatc catatttttc aggggaatct cacattataa ataatgcggc atcgcttggg 420
taaaaacttt tgtgaaagac taaatatgac atgagtctgt ttaaggaagg cgttaaatac 480
gctcagacta cctctggcga attagattta tatttacatg cccctgttga taaggcctta 540
tcacaccacg agcaccttca cttaataaca gtgttaagcg gggcggtatt tcttttccac 600
tcacaccggc cagcgccatg cctttctatg tctcacgcac aagcatccct ctacgtcatc 660
cacgcccgcc tccacactcc ccccgctccg caccgttccc acatagtcgc caccgccatg 720
tccccgctcc cgccccc 737

<210> 1010
<211> 79
<212> PRT
<213> Homo sapiens

<400> 1010
Met Pro Val Trp Leu Gly Gly Thr Phe Ala Pro Leu Cys Leu Ala Cys
1 5 10 15
Arg Ile Ser Asp Asp Phe Gly Glu Cys Cys Cys Ala Pro Tyr Leu Pro
20 25 30

```
Gly Gly Leu His Ser Ile Arg Thr Gly Met Arg Glu Arg Tyr His Ile
        35                  40                  45
Gln Gly Ser Val Gly His Asp Trp Ala Ala Leu Thr Phe Trp Leu Pro
    50                  55                  60
Cys Ala Leu Cys Gln Met Ala Arg Glu Leu Lys Ile Arg Glu  *
65                  70                  75          78
```

<210> 1011
<211> 83
<212> PRT
<213> Homo sapiens

<400> 1011

```
Met Ser Leu Pro Trp Thr Phe Thr Val Leu Ile Leu Ala Pro Ser Leu
1               5                   10                  15
Ser Gly Ser Leu Ser Gly Lys Ser Ser Thr Cys Ala Pro Ala Pro Ser
            20                  25                  30
Ala Pro Gly Ser Arg Ser Ser Gly Pro Arg Arg Asn His His Trp Ile
        35                  40                  45
Ser Arg Tyr Thr Glu Ala Glu Pro Leu Trp Lys Ala Gln Asp Ile Ser
    50                  55                  60
Thr Phe Cys Pro Ser Val Ala Val Thr Phe Arg Gly Asn Ser Val Asn
65                  70                  75                  80
Phe Ala  *
    82
```

<210> 1012
<211> 131
<212> PRT
<213> Homo sapiens

<400> 1012

```
Met Ala Ser Glu Val Val Cys Gly Leu Ile Phe Arg Leu Leu Leu Pro
1               5                   10                  15
Ile Cys Leu Ala Val Ala Cys Ala Phe Arg Tyr Asn Gly Leu Ser Phe
            20                  25                  30
Val Tyr Leu Ile Tyr Leu Leu Leu Ile Pro Leu Phe Ser Glu Pro Thr
        35                  40                  45
Lys Thr Thr Met Gln Gly His Thr Gly Arg Leu Leu Lys Ser Leu Cys
    50                  55                  60
Phe Ile Ser Leu Ser Phe Leu Leu Leu His Ile Ile Phe His Ile Thr
65                  70                  75                  80
Leu Val Ser Leu Glu Ala Gln His Arg Ile Ala Pro Gly Tyr Asn Cys
                85                  90                  95
Ser Thr Trp Glu Lys Thr Phe Arg Gln Ile Gly Phe Glu Ser Leu Lys
            100                 105                 110
Gly Ala Asp Ala Gly Asn Gly Ile Arg Val Leu Val Pro Asp Ile Gly
        115                 120                 125
Met Val Ile
    130 131
```

<210> 1013
<211> 231
<212> PRT
<213> Homo sapiens

<400> 1013
Met Ile Gly Thr Ile Phe Leu Trp Ile Phe Trp Pro Ser Phe Asn Ala
1 5 10 15
Ala Leu Thr Ala Leu Gly Ala Gly Gln His Arg Thr Ala Leu Asn Thr
20 25 30
Tyr Tyr Ser Leu Ala Ala Ser Thr Leu Gly Thr Phe Ala Leu Ser Ala
35 40 45
Leu Val Gly Glu Asp Gly Arg Leu Asp Met Val His Ile Gln Asn Ala
50 55 60
Ala Leu Ala Gly Gly Val Val Val Gly Thr Ser Ser Glu Met Met Leu
65 70 75 80
Thr Pro Phe Gly Ala Leu Ala Ala Gly Phe Leu Ala Gly Thr Val Ser
85 90 95
Thr Leu Gly Tyr Lys Phe Phe Thr Pro Ile Leu Glu Ser Lys Phe Lys
100 105 110
Val Gln Asp Thr Cys Gly Val His Asn Leu His Gly Met Pro Gly Val
115 120 125
Leu Gly Ala Leu Leu Gly Val Leu Val Ala Gly Leu Ala Thr His Glu
130 135 140
Ala Tyr Gly Asp Gly Leu Glu Ser Val Phe Pro Leu Ile Ala Glu Gly
145 150 155 160
Gln Arg Ser Ala Thr Ser Gln Ala Met His Gln Leu Phe Gly Leu Phe
165 170 175
Val Thr Leu Met Phe Ala Ser Val Gly Gly Gly Leu Gly Gly Ile Ile
180 185 190
Leu Val Leu Cys Leu Leu Asp Pro Cys Ala Leu Trp His Trp Val Ala
195 200 205
Pro Ser Ser Met Val Gly Gly Arg Glu Ala Ser Gln Ile Leu Pro Tyr
210 215 220
His His Gln Gly Ser Cys *
225 230

<210> 1014
<211> 60
<212> PRT
<213> Homo sapiens

<400> 1014
Met Cys Glu Ile Ala Asp Leu Trp Ile Gly Leu Leu Trp Leu Phe Phe
1 5 10 15
Val Ile Tyr Cys Phe Ser Phe Asn Ser Leu Thr Thr Val Cys Arg Ala
20 25 30
Ala Val Val Phe Trp Arg Ser Ala Pro Asp Pro Gly Ala Leu Gly Phe
35 40 45
Phe Ser Ile Trp Lys Tyr His Gln Leu Arg Leu *
50 55 59

<210> 1015

<211> 112
<212> PRT
<213> Homo sapiens

<400> 1015
Met Met Thr Val Tyr Pro Leu Leu Gly Tyr Leu Ala Arg Val Gln Leu
1 5 10 15
Leu Gly His Ile Phe Gly Asp Ile Tyr Pro Ser Ile Phe His Val Leu
20 25 30
Ile Leu Asn Leu Ile Ile Val Gly Ala Gly Val Ile Met Ala Cys Phe
35 40 45
Tyr Pro Asn Ile Gly Gly Ile Ile Arg Tyr Ser Gly Ala Ala Cys Gly
50 55 60
Leu Ala Phe Val Phe Ile Tyr Pro Ser Leu Ile Tyr Ile Ile Ser Leu
65 70 75 80
His Gln Glu Glu Arg Leu Thr Trp Pro Lys Leu Ile Phe His Val Phe
85 90 95
Ile Ile Ile Leu Gly Val Ala Asn Leu Ile Val Gln Phe Phe Met *
100 105 110 111

<210> 1016
<211> 68
<212> PRT
<213> Homo sapiens

<400> 1016
Met Ala Lys Tyr Ala Ser Met Thr Phe Lys Leu Phe Ser Leu Cys Val
1 5 10 15
Cys Met Tyr Ile His Ala Cys Thr His Thr His Ile Ser His Thr Asp
20 25 30
Ile Asp Ile Lys Gln Phe Tyr Ala Gln Glu Tyr Gln Gly Gln Pro Lys
35 40 45
Asp Lys Thr Asn Arg Ser Val Ile Tyr Cys Val Phe Asn Phe Ser Thr
50 55 60
Tyr Phe Tyr *
65 67

<210> 1017
<211> 51
<212> PRT
<213> Homo sapiens

<400> 1017
Met Arg Leu Leu Phe Ser Cys Arg Gly Arg Gly Met Phe Leu Phe Arg
1 5 10 15
Arg Arg Met Leu Pro Ser Arg Asp Arg Tyr Tyr Lys Asp Val Glu Leu
20 25 30
Ile Phe Asn Tyr Leu Gly Phe Leu Ile Val Ser Gly Leu Leu Asp Leu
35 40 45
Ile Phe *
50

<210> 1018
<211> 127
<212> PRT
<213> Homo sapiens

<400> 1018
```
Met Leu Arg Phe Tyr Leu Ile Ala Gly Gly Ile Pro Leu Ile Ile Cys
 1              5                   10                  15
Gly Ile Thr Ala Ala Val Asn Ile His Asn Tyr Arg Asp His Ser Pro
            20                  25                  30
Tyr Cys Trp Leu Val Trp Arg Pro Ser Leu Gly Ala Phe Tyr Ile Pro
        35                  40                  45
Val Ala Leu Ile Leu Leu Ile Thr Trp Ile Tyr Phe Leu Cys Ala Gly
    50                  55                  60
Leu Arg Leu Arg Gly Pro Leu Ala Gln Asn Pro Lys Ala Gly Asn Ser
65                  70                  75                  80
Arg Ala Ser Leu Glu Ala Gly Glu Glu Leu Arg Gly Ser Thr Arg Leu
                85                  90                  95
Arg Gly Ser Gly Pro Leu Leu Ser Asp Ser Gly Ser Leu Leu Ala Thr
            100                 105                 110
Gly Ser Ala Arg Val Gly Thr Pro Gly Pro Pro Glu Asp Gly Asp
        115                 120                 125     127
```

<210> 1019
<211> 188
<212> PRT
<213> Homo sapiens

<400> 1019
```
Met Gly Ser Ser Arg Leu Ala Ala Leu Leu Leu Pro Leu Leu Leu Ile
 1              5                   10                  15
Val Ile Asp Leu Ser Asp Ser Ala Gly Ile Gly Phe Arg His Leu Pro
            20                  25                  30
His Trp Asn Thr Arg Cys Pro Leu Ala Ser His Thr Asp Asp Ser Phe
        35                  40                  45
Thr Gly Ser Ser Ala Tyr Ile Pro Cys Arg Thr Trp Trp Ala Leu Phe
    50                  55                  60
Ser Thr Lys Pro Trp Cys Val Arg Val Trp His Cys Ser Arg Cys Leu
65                  70                  75                  80
Cys Gln His Leu Leu Ser Gly Gly Ser Gly Leu Gln Arg Gly Leu Phe
                85                  90                  95
His Leu Leu Val Gln Lys Ser Lys Lys Ser Ser Thr Phe Lys Phe Tyr
            100                 105                 110
Arg Arg His Lys Met Pro Ala Pro Ala Gln Arg Lys Leu Leu Pro Arg
        115                 120                 125
Arg His Leu Ser Glu Lys Ser His His Ile Ser Ile Pro Ser Pro Asp
    130                 135                 140
Ile Ser His Lys Gly Leu Arg Ser Lys Arg Thr Pro Pro Phe Gly Ser
145                 150                 155                 160
Arg Asp Met Gly Lys Ala Phe Pro Lys Trp Asp Ser Pro Thr Pro Gly
                165                 170                 175
Gly Asp Arg Pro Ser Ser Phe Glu Leu Leu Pro  *
            180                 185     187
```

<210> 1020
<211> 65
<212> PRT
<213> Homo sapiens

<400> 1020
Met Ile Leu Leu Cys Pro Gly Leu Thr Asp Leu Ser Val Phe Leu Phe
1 5 10 15
Ser Leu Thr Ile Gly His Phe Ser Arg Val Arg Gly Gln Thr Ile Thr
20 25 30
Ala Cys Pro Ser Ser Arg Ile Pro Ala Gly Phe Gln Asp Ile Val Gln
35 40 45
Gly Ser Ala Asn Ser Gly Pro Arg Ala Leu Ala Arg Cys Pro Cys Leu
50 55 60 64
*

<210> 1021
<211> 136
<212> PRT
<213> Homo sapiens

<400> 1021
Met Pro Gly Phe Lys Phe Cys Ser Ser Leu Arg Phe Leu Tyr Leu Ile
1 5 10 15
Asn Phe Pro Ile Gly Lys Phe Val Cys Leu Ala Ile Leu Leu Pro His
20 25 30
Phe Pro Leu Leu Ser Cys Cys Pro Leu Gln Asp His Leu Asp Phe Pro
35 40 45
Gly Lys Glu Ser Arg Tyr Ser Gly Ser Cys Trp Leu Pro Ser Tyr Ser
50 55 60
Leu Ser Val Ala Gly Ser Pro Leu Gly His Leu Pro Asn Thr Tyr Met
65 70 75 80
His Thr Pro Arg Thr Phe Ser Leu Leu Pro Ile Pro His Pro Ser Val
85 90 95
Asn Trp Asp Ser Phe Lys Pro Phe Ser Ile Arg Glu Ala Leu Ala Thr
100 105 110
Val Glu Ser Leu Gly Arg Gln Ala Phe Pro Asn Thr Pro Thr Thr Trp
115 120 125
Ala Phe Thr Leu His Leu Ser *
130 135

<210> 1022
<211> 186
<212> PRT
<213> Homo sapiens

<400> 1022
Met Ala Gly Pro Arg Pro Arg Trp Arg Asp Gln Leu Leu Phe Met Ser

```
  1               5                   10                  15
Ile Ile Val Leu Val Ile Val Val Ile Cys Leu Met Leu Tyr Ala Leu
            20                  25                  30
Leu Trp Glu Ala Gly Asn Leu Thr Asp Leu Pro Asn Leu Arg Ile Gly
        35                  40                  45
Phe Tyr Asn Phe Cys Leu Trp Asn Glu Asp Thr Ser Thr Leu Gln Cys
    50                  55                  60
His Gln Phe Pro Glu Leu Glu Ala Leu Gly Val Pro Arg Val Gly Leu
65                  70                  75                  80
Gly Leu Ala Arg Leu Gly Val Tyr Gly Ser Leu Val Leu Thr Leu Phe
                85                  90                  95
Ala Pro Gln Pro Leu Leu Leu Ala Gln Cys Asn Ser Asp Glu Arg Ala
            100                 105                 110
Trp Arg Leu Ala Val Gly Phe Leu Ala Val Ser Ser Val Leu Leu Ala
        115                 120                 125
Gly Gly Leu Gly Leu Phe Leu Ser Tyr Val Trp Lys Trp Val Arg Leu
    130                 135                 140
Ser Leu Pro Gly Pro Gly Phe Leu Ala Leu Gly Ser Ala Gln Ala Leu
145                 150                 155                 160
Leu Ile Leu Leu Leu Ile Ala Met Ala Val Phe Pro Leu Arg Ala Glu
                165                 170                 175
Arg Ala Glu Ser Lys Leu Glu Ser Cys  *
            180                 185
```

<210> 1023
<211> 186
<212> PRT
<213> Homo sapiens

<400> 1023

```
Met Ala Gly Pro Arg Pro Arg Trp Arg Asp Gln Leu Leu Phe Met Ser
  1               5                   10                  15
Ile Ile Val Leu Val Ile Val Val Ile Cys Leu Met Leu Tyr Ala Leu
            20                  25                  30
Leu Trp Glu Ala Gly Asn Leu Thr Asp Leu Pro Asn Leu Arg Ile Gly
        35                  40                  45
Phe Tyr Asn Phe Cys Leu Trp Asn Glu Asp Thr Ser Thr Leu Gln Cys
    50                  55                  60
His Gln Phe Pro Glu Leu Glu Ala Leu Gly Val Pro Arg Val Gly Leu
65                  70                  75                  80
Gly Leu Ala Arg Leu Gly Val Tyr Gly Ser Leu Val Leu Thr Leu Phe
                85                  90                  95
Ala Pro Gln Pro Leu Leu Leu Ala Gln Cys Asn Ser Asp Glu Arg Ala
            100                 105                 110
Trp Arg Leu Ala Val Gly Phe Leu Ala Val Ser Ser Val Leu Leu Ala
        115                 120                 125
Gly Gly Leu Gly Leu Phe Leu Ser Tyr Val Trp Lys Trp Val Arg Leu
    130                 135                 140
Ser Leu Pro Gly Pro Gly Phe Leu Ala Leu Gly Ser Ala Gln Ala Leu
145                 150                 155                 160
Leu Ile Leu Leu Leu Ile Ala Met Ala Val Phe Pro Leu Arg Ala Glu
                165                 170                 175
Arg Ala Glu Ser Lys Leu Glu Ser Cys  *
            180                 185
```

<210> 1024
<211> 73
<212> PRT
<213> Homo sapiens

<400> 1024
Met Val Cys Leu Val Gly Phe Leu Glu Leu Ile Leu Tyr Val Tyr Arg
1 5 10 15
Phe Arg Gln Ser Leu Ala Leu Ser His Arg Met Glu Cys Asn Gly Thr
20 25 30
Ile Leu Ala His Cys Asn Leu Arg Leu Pro Gly Ser Ser Asp Ser Pro
35 40 45
Thr Ser Ala Ser Arg Val Ala Gly Ile Thr Gly Thr Arg His His Ala
50 55 60
Arg Val Ile Phe Phe Val Phe Leu *
65 70 72

<210> 1025
<211> 67
<212> PRT
<213> Homo sapiens

<400> 1025
Met Phe Tyr Lys Leu Val Leu Trp Phe Trp Trp Cys Leu Thr Thr Arg
1 5 10 15
Gly Asn Leu Leu Cys Leu Ala Cys Ile Phe Ala Thr Leu Ser Leu Glu
20 25 30
Ser Lys Asn Phe Pro Thr Leu Gln Ala Thr Leu Leu Ile Arg Gln His
35 40 45
Phe Ile Tyr Lys Thr Phe Val Trp Pro Thr Val Cys His Asp Leu Cys
50 55 60
Ser Leu *
65 66

<210> 1026
<211> 67
<212> PRT
<213> Homo sapiens

<400> 1026
Met Gln Ala Gly Ser Ala Leu Trp His Leu Trp Ala Glu Gly Arg Cys
1 5 10 15
Trp Leu Trp Ala Gly Phe Gly Asn Phe Gly Glu Arg Pro His Leu Lys
20 25 30
Thr His Thr Asp Tyr Pro Gly Pro Thr Glu Ala Ser Cys Ile Gln Pro
35 40 45
Tyr Phe Pro Ser Arg Ile Met Leu Ser Ala Thr Pro Leu Glu Gly Tyr
50 55 60
Val Phe *
65 66

<210> 1027
<211> 59
<212> PRT
<213> Homo sapiens

<400> 1027
Met Leu Cys Val Trp Ile Lys Val Leu Phe Leu Leu Ile Ala Glu Ser
1 5 10 15
Asn Thr Trp Leu Leu Ser Pro Arg Thr Lys Asp Val Leu Lys Ser Glu
20 25 30
Pro Thr Gln Ile Tyr Pro His Thr Ser Arg Lys Gln Phe Lys Lys Pro
35 40 45
Gln Glu Ser Lys His Ser Phe Ile Gly Tyr *
50 55 58

<210> 1028
<211> 46
<212> PRT
<213> Homo sapiens

<400> 1028
Met Phe Gln Val Gly Gly Arg Val Phe Lys Arg Cys Ile Phe Ser Phe
1 5 10 15
Cys Cys Cys His Phe Ile Gly Leu Gly Leu Gly Val Cys Phe Ser Ser
20 25 30
Leu Asn Gly Thr Arg Met Phe Ala Asp Ser Tyr Ser Val *
35 40 45

<210> 1029
<211> 61
<212> PRT
<213> Homo sapiens

<400> 1029
Met Ala Phe Arg Thr Cys Phe Leu Ser Cys Leu Thr Val Val Lys Val
1 5 10 15
Cys Ser Lys Ala Ser Pro Ser Phe Ser Thr Gln Gln Pro Cys Val Thr
20 25 30
Thr Lys Val Glu Leu Ser Leu Ile Cys Cys Cys Phe Ser Ser Lys Leu
35 40 45
Pro Asn Lys Ala Lys Asn Thr Leu Val Phe Tyr Ser *
50 55 60

<210> 1030
<211> 50
<212> PRT
<213> Homo sapiens

<400> 1030
Met Trp Leu Arg Lys Cys Leu Leu Gly Leu Ser Leu Ile Ser Phe Arg
1 5 10 15
Val Cys Gly Pro Leu Ile Ala Leu Trp Val Val Ser Asp Ser Ser Ile
20 25 30
Arg Arg Leu Asn Pro Leu Val Val Phe Leu Cys Val Cys Ala Glu Leu
35 40 45
Gly *
49

<210> 1031
<211> 152
<212> PRT
<213> Homo sapiens

<400> 1031
Met Ile Val Tyr Trp Val Leu Met Ser Asn Phe Leu Phe Asn Thr Gly
1 5 10 15
Lys Phe Ile Phe Asn Phe Ile His His Ile Asn Asp Thr Asp Thr Ile
20 25 30
Leu Ser Thr Asn Asn Ser Asn Pro Val Ile Cys Pro Ser Ala Gly Ser
35 40 45
Gly Gly His Pro Asp Asn Ser Ser Met Ile Phe Tyr Ala Asn Asp Thr
50 55 60
Gly Ala Gln Gln Phe Glu Lys Trp Trp Asp Lys Ser Arg Thr Val Pro
65 70 75 80
Phe Tyr Leu Val Gly Leu Leu Leu Pro Leu Leu Asn Phe Lys Ser Pro
85 90 95
Ser Phe Phe Ser Lys Phe Asn Ile Leu Gly Ile Asn Asn Gln Val Ile
100 105 110
Leu Pro Gly Val Thr Glu Met Pro Gly Tyr Cys Pro Phe Leu Leu Pro
115 120 125
Val Ser Thr Glu Cys Cys Ala Val Ala Thr Ser Tyr Thr Cys Phe Glu
130 135 140
Glu Lys Asn Ile Gly Gln Cys Cys
145 150 152

<210> 1032
<211> 1764
<212> PRT
<213> Homo sapiens

<400> 1032
Met Pro Ser Arg Leu Lys Ala Leu Gly Thr Leu Val Ser His Val Thr
1 5 10 15
Leu Arg Leu Leu Lys Pro Glu Cys Val Leu Asp Lys Ser Trp Cys Gln
20 25 30
Glu Glu Leu Ser Val Ala Val Lys Arg Ala Val Met Leu Leu His Thr
35 40 45
His Thr Ile Thr Ser Arg Val Gly Lys Gly Glu Pro Gly Ala Ala Pro
50 55 60
Leu Ser Ala Pro Ala Phe Ser Leu Val Phe Pro Phe Leu Lys Met Val

```
65                  70                  75                  80
Leu Thr Glu Met Pro His His Ser Glu Glu Glu Glu Glu Trp Met Ala
                85                  90                  95
Gln Ile Leu Gln Ile Leu Thr Val Gln Ala Gln Leu Arg Ala Ser Pro
            100                 105                 110
Asn Thr Pro Pro Gly Arg Val Asp Glu Asn Gly Pro Glu Leu Leu Pro
        115                 120                 125
Arg Val Ala Met Leu Arg Leu Leu Thr Trp Val Ile Gly Thr Gly Ser
    130                 135                 140
Pro Arg Leu Gln Val Leu Ala Ser Asp Thr Leu Thr Thr Leu Cys Ala
145                 150                 155                 160
Ser Ser Ser Gly Asp Asp Gly Cys Ala Phe Ala Glu Gln Glu Glu Val
                165                 170                 175
Asp Val Leu Leu Cys Ala Leu Gln Ser Pro Cys Ala Ser Val Arg Glu
            180                 185                 190
Thr Val Leu Arg Gly Leu Met Glu Leu His Met Val Leu Pro Ala Pro
        195                 200                 205
Asp Thr Asp Glu Lys Asn Gly Leu Asn Leu Leu Arg Arg Leu Trp Val
    210                 215                 220
Val Lys Phe Asp Lys Glu Glu Glu Ile Arg Lys Leu Ala Glu Arg Leu
225                 230                 235                 240
Trp Ser Met Met Gly Leu Asp Leu Gln Pro Asp Leu Cys Ser Leu Leu
                245                 250                 255
Ile Asp Asp Val Ile Tyr His Glu Ala Ala Val Arg Gln Ala Gly Ala
            260                 265                 270
Glu Ala Leu Ser Gln Ala Val Ala Arg Tyr Gln Arg Gln Ala Ala Glu
        275                 280                 285
Val Met Gly Arg Leu Met Glu Ile Tyr Gln Glu Lys Leu Tyr Arg Pro
    290                 295                 300
Pro Pro Val Leu Asp Ala Leu Gly Arg Val Ile Ser Glu Ser Pro Pro
305                 310                 315                 320
Asp Gln Trp Glu Ala Arg Cys Gly Leu Ala Leu Ala Leu Asn Lys Leu
                325                 330                 335
Ser Gln Tyr Leu Asp Ser Ser Gln Val Lys Pro Leu Phe Gln Phe Phe
            340                 345                 350
Val Pro Asp Ala Leu Asn Asp Arg His Pro Asp Val Arg Lys Cys Met
        355                 360                 365
Leu Asp Ala Ala Leu Ala Thr Leu Asn Thr His Gly Lys Glu Asn Val
    370                 375                 380
Asn Ser Leu Leu Pro Val Phe Glu Glu Phe Leu Lys Asn Ala Pro Asn
385                 390                 395                 400
Asp Ala Ser Tyr Asp Ala Val Arg Gln Ser Val Val Val Leu Met Gly
                405                 410                 415
Ser Leu Ala Lys His Leu Asp Lys Ser Asp Pro Lys Val Lys Pro Ile
            420                 425                 430
Val Ala Lys Leu Ile Ala Ala Leu Ser Thr Pro Ser Gln Gln Val Gln
        435                 440                 445
Glu Ser Val Ala Ser Cys Leu Pro Pro Leu Val Pro Ala Ile Lys Glu
    450                 455                 460
Asp Ala Gly Gly Met Ile Gln Arg Leu Met Gln Gln Leu Leu Glu Ser
465                 470                 475                 480
Asp Lys Tyr Ala Glu Arg Lys Gly Ala Ala Tyr Gly Leu Ala Gly Leu
                485                 490                 495
Val Lys Gly Leu Gly Ile Leu Ser Leu Lys Gln Gln Glu Met Met Ala
            500                 505                 510
Ala Leu Thr Asp Ala Ile Gln Asp Lys Lys Asn Phe Arg Arg Arg Glu
        515                 520                 525
Gly Ala Leu Phe Ala Phe Glu Met Leu Cys Thr Met Leu Gly Lys Leu
    530                 535                 540
```

Phe Glu Pro Tyr Val Val His Val Leu Pro His Leu Leu Leu Cys Phe
545 550 555 560
Gly Asp Gly Asn Gln Tyr Val Arg Glu Ala Ala Asp Asp Cys Ala Lys
565 570 575
Ala Val Met Ser Asn Leu Ser Ala His Gly Val Lys Leu Val Leu Pro
580 585 590
Ser Leu Leu Ala Ala Leu Glu Glu Glu Ser Trp Arg Thr Lys Ala Gly
595 600 605
Ser Val Glu Leu Leu Gly Ala Met Ala Tyr Cys Ala Pro Lys Gln Leu
610 615 620
Ser Ser Cys Leu Pro Asn Ile Val Pro Lys Leu Thr Glu Val Leu Thr
625 630 635 640
Asp Ser His Val Lys Val Gln Lys Ala Gly Gln Gln Ala Leu Arg Gln
645 650 655
Ile Gly Ser Val Ile Arg Asn Pro Glu Ile Leu Ala Ile Ala Pro Val
660 665 670
Leu Leu Asp Ala Leu Thr Asp Pro Ser Arg Lys Thr Gln Lys Cys Leu
675 680 685
Gln Thr Leu Leu Asp Thr Lys Phe Val His Phe Ile Asp Ala Pro Ser
690 695 700
Leu Ala Leu Ile Met Pro Ile Val Gln Arg Ala Phe Gln Asp Arg Ser
705 710 715 720
Thr Asp Thr Arg Lys Met Ala Ala Gln Ile Ile Gly Asn Met Tyr Ser
725 730 735
Leu Thr Asp Gln Lys Asp Leu Ala Pro Tyr Leu Pro Ser Val Thr Pro
740 745 750
Gly Leu Lys Ala Ser Leu Leu Asp Pro Val Pro Glu Val Arg Thr Val
755 760 765
Ser Ala Lys Ala Leu Gly Ala Met Val Lys Gly Met Gly Glu Ser Cys
770 775 780
Phe Glu Asp Leu Leu Pro Trp Leu Met Glu Thr Leu Thr Tyr Glu Gln
785 790 795 800
Ser Ser Val Asp Arg Ser Gly Ala Ala Gln Gly Leu Ala Glu Val Met
805 810 815
Ala Gly Leu Gly Val Glu Lys Leu Glu Lys Leu Met Pro Glu Ile Val
820 825 830
Ala Thr Ala Ser Lys Val Asp Ile Ala Pro His Val Arg Asp Gly Tyr
835 840 845
Ile Met Met Phe Asn Tyr Leu Pro Ile Thr Phe Gly Asp Lys Phe Thr
850 855 860
Pro Tyr Val Gly Pro Ile Ile Pro Cys Ile Leu Lys Ala Leu Ala Asp
865 870 875 880
Glu Asn Glu Phe Val Arg Asp Thr Ala Leu Arg Ala Gly Gln Arg Val
885 890 895
Ile Ser Met Tyr Ala Glu Thr Ala Ile Ala Leu Leu Leu Pro Gln Leu
900 905 910
Glu Gln Gly Leu Phe Asp Asp Leu Trp Arg Ile Arg Phe Ser Ser Val
915 920 925
Gln Leu Leu Gly Asp Leu Leu Phe His Ile Ser Gly Val Thr Gly Lys
930 935 940
Met Thr Thr Glu Thr Ala Ser Glu Asp Asp Asn Phe Gly Thr Ala Gln
945 950 955 960
Ser Asn Lys Ala Ile Ile Thr Ala Leu Gly Val Glu Arg Arg Asn Arg
965 970 975
Val Leu Ala Gly Leu Tyr Met Gly Arg Ser Asp Thr Gln Leu Val Val
980 985 990
Arg Gln Ala Ser Leu His Val Trp Lys Ile Val Val Ser Asn Thr Pro
995 1000 1005
Arg Thr Leu Arg Glu Ile Leu Pro Thr Leu Phe Gly Leu Leu Leu Gly

1010 1015 1020
Phe Leu Ala Ser Thr Cys Ala Asp Lys Arg Thr Ile Ala Ala Arg Thr
1025 1030 1035 1040
Leu Gly Asp Leu Val Arg Lys Leu Gly Glu Lys Ile Leu Pro Glu Ile
1045 1050 1055
Ile Pro Ile Leu Glu Glu Gly Leu Arg Ser Gln Lys Ser Asp Glu Arg
1060 1065 1070
Gln Gly Val Cys Ile Gly Leu Ser Glu Ile Met Lys Ser Thr Ser Arg
1075 1080 1085
Asp Ala Val Leu Tyr Phe Ser Glu Ser Leu Val Pro Thr Ala Arg Lys
1090 1095 1100
Ala Leu Cys Asp Pro Leu Glu Glu Val Arg Glu Ala Ala Ala Lys Thr
1105 1110 1115 1120
Phe Glu Gln Leu His Ser Thr Ile Gly His Gln Ala Leu Glu Asp Ile
1125 1130 1135
Leu Pro Phe Leu Leu Lys Gln Leu Asp Asp Glu Glu Val Ser Glu Phe
1140 1145 1150
Ala Leu Asp Gly Leu Lys Gln Val Met Ala Ile Lys Ser Arg Val Val
1155 1160 1165
Leu Pro Tyr Leu Val Pro Lys Leu Thr Thr Pro Pro Val Asn Thr Arg
1170 1175 1180
Val Leu Ala Phe Leu Ser Ser Val Ala Gly Asp Ala Leu Thr Arg His
1185 1190 1195 1200
Leu Gly Val Ile Leu Pro Ala Val Met Leu Ala Leu Lys Glu Lys Leu
1205 1210 1215
Gly Thr Pro Asp Glu Gln Leu Glu Met Ala Asn Cys Gln Ala Val Ile
1220 1225 1230
Leu Ser Val Glu Asp Asp Thr Gly His Arg Ile Ile Ile Glu Asp Leu
1235 1240 1245
Leu Glu Ala Thr Arg Ser Pro Glu Val Gly Met Arg Gln Ala Ala Ala
1250 1255 1260
Ile Ile Leu Asn Ile Tyr Cys Ser Arg Ser Lys Ala Asp Tyr Thr Ser
1265 1270 1275 1280
His Leu Arg Ser Leu Val Ser Gly Leu Ile Arg Leu Phe Asn Asp Ser
1285 1290 1295
Ser Pro Val Val Leu Glu Glu Ser Trp Asp Ala Leu Asn Ala Ile Thr
1300 1305 1310
Lys Lys Leu Asp Ala Gly Asn Gln Leu Ala Leu Ile Glu Glu Leu His
1315 1320 1325
Lys Glu Ile Arg Leu Ile Gly Asn Glu Ser Lys Gly Glu His Val Pro
1330 1335 1340
Gly Phe Cys Leu Pro Lys Lys Gly Val Thr Ser Ile Leu Pro Val Leu
1345 1350 1355 1360
Arg Glu Gly Val Leu Thr Gly Ser Pro Glu Gln Lys Glu Glu Ala Ala
1365 1370 1375
Lys Ala Leu Gly Leu Val Ile Arg Leu Thr Ser Ala Asp Ala Leu Arg
1380 1385 1390
Pro Ser Val Val Ser Ile Thr Gly Pro Leu Ile Arg Ile Leu Gly Asp
1395 1400 1405
Arg Phe Ser Trp Asn Val Lys Ala Ala Leu Leu Glu Thr Leu Ser Leu
1410 1415 1420
Leu Leu Ala Lys Val Gly Ile Ala Leu Lys Pro Phe Leu Pro Gln Leu
1425 1430 1435 1440
Gln Thr Thr Phe Thr Lys Ala Leu Gln Asp Ser Asn Arg Gly Val Arg
1445 1450 1455
Leu Lys Ala Ala Asp Ala Leu Gly Lys Leu Ile Ser Ile His Ile Lys
1460 1465 1470
Val Asp Pro Leu Phe Thr Glu Leu Leu Asn Gly Ile Arg Ala Met Glu
1475 1480 1485

Asp Pro Gly Val Arg Asp Thr Met Leu Gln Ala Leu Arg Phe Val Ile
1490 1495 1500
Gln Gly Ala Gly Ala Lys Val Asp Ala Val Ile Arg Lys Asn Ile Val
1505 1510 1515 1520
Ser Leu Leu Leu Ser Met Leu Gly His Asp Glu Asp Asn Thr Arg Ile
1525 1530 1535
Ser Ser Ala Gly Cys Leu Gly Glu Leu Cys Ala Phe Leu Thr Glu Glu
1540 1545 1550
Glu Leu Ser Ala Val Leu Gln Gln Cys Leu Leu Ala Asp Val Ser Gly
1555 1560 1565
Ile Asp Trp Met Val Arg His Gly Arg Ser Leu Ala Leu Ser Val Ala
1570 1575 1580
Val Asn Val Ala Pro Gly Arg Leu Cys Ala Gly Arg Tyr Ser Ser Asp
1585 1590 1595 1600
Val Gln Glu Met Ile Leu Ser Ser Ala Thr Ala Asp Arg Ile Pro Ile
1605 1610 1615
Ala Val Ser Gly Val Arg Gly Met Gly Phe Leu Met Arg His His Ile
1620 1625 1630
Glu Thr Gly Gly Gly Gln Leu Pro Ala Lys Leu Ser Ser Leu Phe Val
1635 1640 1645
Lys Cys Leu Gln Asn Pro Ser Ser Asp Ile Arg Leu Val Ala Glu Lys
1650 1655 1660
Met Ile Trp Trp Ala Asn Lys Asp Pro Leu Pro Pro Leu Asp Pro Gln
1665 1670 1675 1680
Ala Ile Lys Pro Ile Leu Lys Ala Leu Leu Asp Asn Thr Lys Asp Lys
1685 1690 1695
Asn Thr Val Val Arg Ala Tyr Ser Asp Gln Ala Ile Val Asn Leu Leu
1700 1705 1710
Lys Met Arg Gln Gly Glu Glu Val Phe Gln Ser Leu Ser Lys Ile Leu
1715 1720 1725
Asp Val Ala Ser Leu Glu Val Leu Asn Glu Val Asn Arg Arg Ser Leu
1730 1735 1740
Lys Lys Leu Ala Ser Gln Ala Asp Ser Thr Glu Gln Val Asp Asp Thr
1745 1750 1755 1760
Ile Leu Thr *
1763

<210> 1033
<211> 151
<212> PRT
<213> Homo sapiens

<400> 1033
Met Asn Arg Arg Ala Ser Gln Met Leu Leu Met Phe Leu Leu Ala Ile
1 5 10 15
Cys Leu Leu Ala Ile Ile Phe Val Pro Gln Glu Met Gln Met Leu Arg
20 25 30
Glu Val Leu Ala Thr Leu Gly Leu Gly Ala Ser Ala Leu Ala Asn Thr
35 40 45
Leu Ala Phe Ala His Gly Asn Glu Val Ile Pro Thr Ile Ile Arg Ala
50 55 60
Arg Ala Met Gly Ile Asn Ala Thr Phe Ala Asn Ile Ala Gly Ala Leu
65 70 75 80
Ala Pro Leu Met Met Ile Leu Ser Val Tyr Ser Pro Pro Leu Pro Trp
85 90 95
Ile Ile Tyr Gly Val Phe Pro Phe Ile Ser Gly Phe Ala Phe Leu Leu

```
            100                 105                 110
Leu Pro Glu Thr Arg Asn Lys Pro Leu Phe Asp Thr Ile Gln Asp Glu
        115                 120                 125
Lys Asn Glu Arg Lys Asp Pro Arg Glu Pro Lys Gln Glu Asp Pro Arg
    130                 135                 140
Val Glu Val Thr Gln Phe  *
145                 150
```

```
     <210> 1034
     <211> 149
     <212> PRT
     <213> Homo sapiens


     <400> 1034
Met Ala Leu Leu Leu Pro Arg Trp Phe Arg Glu Ala Pro Val Leu Phe
  1               5                  10                  15
Ser Thr Gly Trp Ser Pro Leu Asp Val Leu Leu His Ser Leu Leu Thr
             20                  25                  30
Gln Pro Ile Phe Leu Ala Gly Leu Ser Gly Phe Leu Leu Glu Asn Thr
         35                  40                  45
Ile Pro Gly Thr Gln Leu Glu Arg Gly Leu Gly Gln Gly Leu Pro Ser
     50                  55                  60
Pro Phe Thr Ala Gln Glu Ala Arg Met Pro Gln Lys Pro Arg Glu Lys
 65                  70                  75                  80
Ala Ala Gln Val Tyr Arg Leu Pro Phe Pro Ile Gln Asn Leu Cys Pro
                 85                  90                  95
Cys Ile Pro Gln Pro Leu His Cys Leu Cys Pro Leu Pro Glu Asp Pro
            100                 105                 110
Gly Asp Glu Glu Gly Gly Ser Ser Glu Pro Glu Glu Met Ala Asp Leu
        115                 120                 125
Leu Pro Gly Ser Gly Glu Pro Cys Pro Glu Ser Thr Arg Glu Gly Val
    130                 135                 140
Arg Ser Gln Lys  *
145         148
```

```
     <210> 1035
     <211> 88
     <212> PRT
     <213> Homo sapiens


     <400> 1035
Met Gly Ile Ala Leu Leu Gln Ile Phe Gly Ile Cys Leu Ala Gln Asn
  1               5                  10                  15
Leu Val Ser Asp Ile Lys Ala Val Lys Ala Asn Trp Ser Lys Trp Asn
             20                  25                  30
Asp Asp Phe Glu Asn His Trp Leu Thr Pro Thr Ile Ser Glu Val Leu
         35                  40                  45
Ser Thr Ala Gly Pro Gln Gln Asn Ser Leu Thr Gly Ala Pro Gly Pro
     50                  55                  60
Ala Pro Pro Ser Arg His Val Phe Phe Gly Leu Gly Gly Leu Tyr Pro
 65                  70                  75                  80
Glu Pro Thr Phe Lys Asn Trp  *
                 85      87
```

<210> 1036
<211> 96
<212> PRT
<213> Homo sapiens

<400> 1036
Met Val Val Leu Ile Pro Val Ser Trp Val Ala Asn Ala Ile Ile Arg
1 5 10 15
Asp Phe Tyr Asn Ser Ile Val Asn Val Ala Gln Lys Arg Glu Leu Gly
20 25 30
Glu Ala Leu Tyr Leu Gly Trp Thr Thr Ala Leu Val Leu Ile Val Gly
35 40 45
Gly Ala Leu Phe Cys Cys Val Phe Cys Cys Asn Glu Lys Ser Ser Ser
50 55 60
Tyr Arg Tyr Ser Ile Pro Ser His Arg Thr Thr Gln Lys Ser Tyr His
65 70 75 80
Thr Gly Lys Lys Ser Pro Ser Val Tyr Ser Arg Ser Gln Tyr Val *
85 90 95

<210> 1037
<211> 139
<212> PRT
<213> Homo sapiens

<400> 1037
Met Ala Leu Ser Trp Met Thr Ile Val Val Pro Leu Leu Thr Phe Glu
1 5 10 15
Ile Leu Leu Val His Lys Leu Asp Gly His Asn Ala Phe Ser Cys Ile
20 25 30
Pro Ile Phe Val Pro Leu Trp Leu Ser Leu Ile Thr Leu Met Ala Thr
35 40 45
Thr Phe Gly Gln Lys Gly Gly Asn His Trp Trp Phe Gly Ile Arg Lys
50 55 60
Asp Phe Cys Gln Phe Leu Leu Glu Ile Phe Pro Phe Leu Arg Glu Tyr
65 70 75 80
Gly Asn Ile Ser Tyr Asp Leu His His Glu Asp Asn Glu Glu Thr Glu
85 90 95
Glu Thr Pro Val Pro Glu Pro Pro Lys Ile Ala Pro Met Phe Arg Lys
100 105 110
Lys Ala Arg Val Val Ile Thr Gln Ser Pro Gly Lys Tyr Val Leu Pro
115 120 125
Pro Pro Lys Leu Asn Ile Glu Met Pro Asp *
130 135 138

<210> 1038
<211> 64
<212> PRT
<213> Homo sapiens

<400> 1038
Met Val Leu Ser Gly Ile His Trp Tyr Ser Val Leu Leu Leu Ala Val
1 5 10 15
Glu Phe Cys Arg Tyr Cys Pro Leu Arg Tyr Arg Cys Ser Thr Phe Ser
20 25 30
Ser Trp Ala Arg Val Ser Ser Thr Pro Gln Ala Ser Ser Pro Val Ala
35 40 45
Leu Thr Met Leu Ser Ser Arg Gly Arg Ser Glu Gly Gly Ala Leu *
50 55 60 63

<210> 1039
<211> 286
<212> PRT
<213> Homo sapiens

<400> 1039
Met Met Leu Gly Pro Val Thr Leu His Leu Val Gly His Leu Leu Ala
1 5 10 15
Phe Leu Asp Leu Leu Cys Pro Arg Gly Pro Ile His Ser Ile Leu Pro
20 25 30
Met Thr Phe Glu Ala Val Lys Gln Asp His Gly Phe Met Leu Tyr Arg
35 40 45
Thr Tyr Met Thr His Thr Ile Phe Glu Pro Thr Pro Phe Trp Val Pro
50 55 60
Asn Asn Gly Val His Asp Arg Ala Tyr Val Met Val Asp Gly Val Phe
65 70 75 80
Gln Gly Val Val Glu Arg Asn Met Arg Asp Lys Leu Phe Leu Thr Gly
85 90 95
Lys Leu Gly Ser Lys Leu Asp Ile Leu Val Glu Asn Met Gly Arg Leu
100 105 110
Ser Phe Gly Ser Asn Ser Ser Asp Phe Lys Gly Leu Leu Lys Pro Pro
115 120 125
Ile Leu Gly Gln Thr Ile Leu Thr Gln Trp Met Met Phe Pro Leu Lys
130 135 140
Ile Asp Asn Leu Val Lys Trp Trp Phe Pro Leu Gln Leu Pro Lys Trp
145 150 155 160
Pro Tyr Pro Gln Ala Pro Ser Gly Pro Thr Phe Tyr Ser Lys Thr Phe
165 170 175
Pro Ile Leu Gly Ser Val Gly Asp Thr Phe Leu Tyr Leu Pro Gly Trp
180 185 190
Thr Lys Gly Gln Val Trp Ile Asn Gly Phe Asn Leu Gly Arg Tyr Trp
195 200 205
Thr Lys Gln Gly Pro Gln Gln Thr Leu Tyr Val Pro Arg Phe Leu Leu
210 215 220
Phe Pro Arg Gly Ala Leu Asn Lys Ile Thr Leu Leu Glu Leu Glu Asp
225 230 235 240
Val Pro Leu Gln Pro Gln Val Gln Phe Leu Asp Lys Pro Ile Leu Asn
245 250 255
Ser Thr Ser Thr Leu His Arg Thr His Ile Asn Ser Leu Ser Ala Asp
260 265 270
Thr Leu Ser Ala Ser Glu Pro Met Glu Leu Ser Gly His *
275 280 285

<210> 1040

<211> 96
<212> PRT
<213> Homo sapiens

<400> 1040
Met His Ala His Ser Ala Ser Leu Trp Val Ala Phe Phe Tyr Arg Ser
1 5 10 15
Pro Phe Leu Phe Phe Thr Thr Gly Pro Pro Pro Pro Thr Ser Ser Ser
20 25 30
Pro Ala Gly Leu Pro Leu Leu Glu Ser Thr Val Asp Ala Ser Arg Pro
35 40 45
Asn Trp Leu Pro Leu Leu Leu Ser Pro Pro Leu Pro Phe Leu Ser Ile
50 55 60
Glu Cys Thr Leu Tyr Asn Phe Ser Gly Ile Val Ile Glu Asn Lys Ile
65 70 75 80
Phe Thr Ile Ile Thr Gly Phe Phe Gln Val Thr Ser Cys Arg Leu *
85 90 95

<210> 1041
<211> 64
<212> PRT
<213> Homo sapiens

<400> 1041
Met Ser Asp Ile Ser Pro Leu Leu Tyr Glu Ile Trp Leu Gly Asp Thr
1 5 10 15
Ser Ala Gly Phe Phe Thr Phe Cys Val Thr Val Leu His Val Leu Leu
20 25 30
Leu Leu Ser Ser Val Leu His Phe Leu Cys Pro Arg Asp Thr Ser Val
35 40 45
Ile Ser Pro Phe Ile Pro Pro Leu Thr Pro Pro Gln Ser Arg Leu *
50 55 60 63

<210> 1042
<211> 415
<212> PRT
<213> Homo sapiens

<400> 1042
Met Asn Glu Thr Gly Val Ile Val Trp Tyr Leu Ala Leu Cys Leu Leu
1 5 10 15
Leu Ala Trp Leu Ile Val Gly Ala Ala Leu Phe Lys Gly Ile Lys Ser
20 25 30
Ser Gly Lys Val Val Tyr Phe Thr Ala Leu Phe Pro Tyr Val Val Leu
35 40 45
Leu Ile Leu Leu Val Arg Gly Ala Thr Leu Glu Gly Ala Ser Lys Gly
50 55 60
Ile Ser Tyr Tyr Ile Gly Ala Gln Ser Asn Phe Thr Lys Leu Lys Glu
65 70 75 80
Ala Glu Val Trp Lys Asp Ala Ala Thr Gln Ile Phe Tyr Ser Leu Ser
85 90 95
Val Ala Trp Gly Gly Leu Val Ala Leu Ser Ser Tyr Asn Lys Phe Lys

```
                100                 105                 110
Asn Asn Cys Phe Ser Asp Ala Ile Val Val Cys Leu Thr Asn Cys Leu
            115                 120                 125
Thr Ser Val Phe Ala Gly Phe Ala Ile Phe Ser Ile Leu Gly His Met
        130                 135                 140
Ala His Ile Ser Gly Lys Glu Val Ser Gln Val Val Lys Ser Gly Phe
145                 150                 155                 160
Asp Leu Ala Phe Ile Ala Tyr Pro Glu Ala Leu Ala Gln Leu Pro Gly
                165                 170                 175
Gly Pro Phe Trp Ser Ile Leu Phe Phe Phe Met Leu Leu Thr Leu Gly
            180                 185                 190
Leu Asp Ser Gln Phe Ala Ser Ile Glu Thr Ile Thr Thr Thr Ile Gln
        195                 200                 205
Asp Leu Phe Pro Lys Val Met Lys Lys Met Arg Val Pro Ile Thr Leu
    210                 215                 220
Gly Cys Cys Leu Val Leu Phe Leu Leu Gly Leu Val Cys Val Thr Gln
225                 230                 235                 240
Ala Gly Ile Tyr Trp Val His Leu Ile Asp His Phe Cys Ala Gly Trp
                245                 250                 255
Gly Ile Leu Ile Ala Ala Ile Leu Glu Leu Val Gly Ile Ile Trp Ile
            260                 265                 270
Tyr Gly Gly Asn Arg Phe Ile Glu Asp Thr Glu Met Met Ile Gly Ala
        275                 280                 285
Lys Arg Trp Ile Phe Trp Leu Trp Trp Arg Ala Cys Trp Phe Val Ile
    290                 295                 300
Thr Pro Ile Leu Leu Ile Ala Ile Phe Ile Trp Ser Leu Val Gln Phe
305                 310                 315                 320
His Arg Pro Asn Tyr Gly Ala Ile Pro Tyr Pro Asp Trp Gly Val Ala
                325                 330                 335
Leu Gly Trp Cys Met Ile Val Phe Cys Ile Ile Trp Ile Pro Ile Met
            340                 345                 350
Ala Ile Ile Lys Ile Ile Gln Ala Lys Gly Asn Ile Phe Gln Arg Leu
        355                 360                 365
Ile Ser Cys Cys Arg Pro Ala Ser Asn Trp Gly Pro Tyr Leu Glu Gln
    370                 375                 380
His Arg Gly Glu Arg Tyr Lys Asp Met Val Asp Pro Lys Lys Glu Ala
385                 390                 395                 400
Asp His Glu Ile Pro Thr Val Ser Gly Ser Arg Lys Pro Glu  *
                405                 410                 414
```

```
<210> 1043
<211> 48
<212> PRT
<213> Homo sapiens
```

```
<400> 1043
Met Pro Thr Leu Gly Asp Ala Leu Ile Leu Tyr Leu His Leu Val Leu
 1               5                  10                  15
Gly Val Ala Gly Val Leu Gln Pro Pro Gly Pro Arg Pro Ser Gln Ala
            20                  25                  30
Leu Gly Pro Thr Gly Asp Arg Ala Pro Gly Lys Trp Asn Arg Ser  *
        35                  40                  45          47
```

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<210> 1044
```

<211> 146
<212> PRT
<213> Homo sapiens

<400> 1044
Met Leu Phe Ser Ser Met Thr Leu Arg Leu Ser Arg Cys Ser Cys Ser
1 5 10 15
Ile Leu Leu Phe Trp Ala Ser Ala Ala Cys Met Phe Pro Ser Ser Arg
20 25 30
Tyr Leu Trp Ser Gly Arg Ser Leu Val Ser Val Glu Gly Ser Asp Arg
35 40 45
Phe Ser Ser Ala Val Ser Ser Phe Ser Ser Lys Ala Asn Trp Val Lys
50 55 60
Pro Lys Phe Arg Ser Trp Ser Gly Gly Ile Glu Leu Gly Phe Gln Met
65 70 75 80
His Trp Pro Pro Gly Val Gly Pro Arg Tyr Ser Pro Ser Cys His Phe
85 90 95
Pro Lys Ser Arg Trp Arg Thr Arg Pro Leu Arg Leu Ser Thr Ala Pro
100 105 110
Cys Thr Ser Trp Thr Leu Glu Leu Gln Tyr Leu Ala Leu Gln Lys Val
115 120 125
Ile Leu Gln Trp Gln Glu Leu Ser Cys Val Phe Arg Met Ser Thr Ser
130 135 140
Pro *
145

<210> 1045
<211> 53
<212> PRT
<213> Homo sapiens

<400> 1045
Met Ala Leu Phe Cys Leu Val Tyr Gln Ile Ile Phe Leu Ile Gln His
1 5 10 15
Thr His Phe Ser Leu Ala Lys Leu Leu Ile Met Ala Leu Asn Thr Leu
20 25 30
Thr Tyr Cys Val Leu Val Gln Ser Asn Asn Thr Gln Ser Thr Leu Arg
35 40 45
Lys Ser Ala Ser *
50 52

<210> 1046
<211> 407
<212> PRT
<213> Homo sapiens

<400> 1046
Met Gly Pro Ser Thr Pro Leu Leu Ile Leu Phe Leu Leu Ser Trp Ser
1 5 10 15
Gly Pro Leu Gln Gly Gln Gln His His Leu Val Glu Tyr Met Glu Arg
20 25 30
Arg Leu Ala Ala Leu Glu Glu Arg Leu Ala Gln Cys Gln Asp Gln Ser

```
        35                  40                  45
Ser Arg His Ala Ala Glu Leu Arg Asp Phe Lys Asn Lys Met Leu Pro
    50                  55                  60
Leu Leu Glu Val Ala Glu Lys Glu Arg Glu Ala Leu Arg Thr Glu Ala
65                  70                  75                  80
Asp Thr Ile Ser Gly Arg Val Asp Arg Leu Glu Arg Glu Val Asp Tyr
                85                  90                  95
Leu Glu Thr Gln Asn Pro Ala Leu Pro Cys Val Glu Phe Asp Glu Lys
            100                 105                 110
Val Thr Gly Gly Pro Gly Thr Lys Gly Lys Gly Arg Arg Asn Glu Lys
        115                 120                 125
Tyr Asp Met Val Thr Asp Cys Gly Tyr Thr Ile Ser Gln Val Arg Ser
    130                 135                 140
Met Lys Ile Leu Lys Arg Phe Gly Gly Pro Ala Gly Leu Trp Thr Lys
145                 150                 155                 160
Asp Pro Leu Gly Gln Thr Glu Lys Ile Tyr Val Leu Asp Gly Thr Gln
                165                 170                 175
Asn Asp Thr Ala Phe Val Phe Pro Arg Leu Arg Asp Phe Thr Leu Ala
            180                 185                 190
Met Ala Ala Arg Lys Ala Ser Arg Val Arg Val Pro Phe Pro Trp Val
        195                 200                 205
Gly Thr Gly Gln Leu Val Tyr Gly Gly Phe Leu Tyr Phe Ala Arg Arg
    210                 215                 220
Pro Pro Gly Arg Pro Gly Gly Gly Gly Glu Met Glu Asn Thr Leu Gln
225                 230                 235                 240
Leu Ile Lys Phe His Leu Ala Asn Arg Thr Val Val Asp Ser Ser Val
                245                 250                 255
Phe Pro Ala Glu Gly Leu Ile Pro Pro Tyr Gly Leu Thr Ala Asp Thr
            260                 265                 270
Tyr Ile Asp Leu Ala Ala Asp Glu Glu Gly Leu Trp Ala Val Tyr Ala
        275                 280                 285
Thr Arg Glu Asp Asp Arg His Leu Cys Leu Ala Lys Leu Asp Pro Gln
    290                 295                 300
Thr Leu Asp Thr Glu Gln Gln Trp Asp Thr Pro Cys Pro Arg Glu Asn
305                 310                 315                 320
Ala Glu Ala Ala Phe Val Ile Cys Gly Thr Leu Tyr Val Val Tyr Asn
                325                 330                 335
Thr Arg Pro Ala Ser Arg Ala Arg Ile Gln Cys Ser Phe Asp Ala Ser
            340                 345                 350
Gly Thr Leu Thr Pro Glu Arg Ala Ala Leu Pro Tyr Phe Pro Arg Arg
        355                 360                 365
Tyr Gly Ala His Ala Ser Leu Arg Tyr Asn Pro Arg Glu Arg Gln Leu
    370                 375                 380
Tyr Ala Trp Asp Asp Gly Tyr Gln Ile Val Tyr Lys Leu Glu Met Arg
385                 390                 395                 400
Lys Lys Glu Glu Glu Val  *
                405 406
```

<210> 1047
<211> 268
<212> PRT
<213> Homo sapiens

<400> 1047

```
Met Ile Gln Lys Ile Leu Phe Lys Asp Leu Phe Arg Phe Leu Leu Val
 1               5                  10                  15
```

Tyr Leu Leu Phe Met Ile Gly Tyr Ala Ser Ala Leu Val Ser Leu Leu
20 25 30
Asn Pro Cys Ala Asn Met Lys Val Cys Asn Glu Asp Gln Thr Asn Cys
35 40 45
Thr Val Pro Thr Tyr Pro Ser Cys Arg Asp Ser Glu Thr Phe Ser Thr
50 55 60
Phe Leu Leu Asp Leu Phe Lys Leu Thr Ile Gly Met Gly Asp Leu Glu
65 70 75 80
Met Leu Ser Ser Thr Lys Tyr Pro Val Val Phe Ile Ile Leu Leu Val
85 90 95
Thr Tyr Ile Ile Leu Thr Phe Val Leu Leu Leu Asn Met Leu Ile Ala
100 105 110
Leu Met Gly Glu Thr Val Gly Gln Val Ser Lys Glu Ser Lys His Ile
115 120 125
Trp Lys Leu Gln Trp Ala Thr Thr Ile Leu Asp Ile Glu Arg Ser Phe
130 135 140
Pro Val Phe Leu Arg Lys Ala Phe Arg Ser Gly Glu Met Val Thr Val
145 150 155 160
Gly Lys Ser Ser Asp Gly Thr Pro Asp Arg Arg Trp Cys Phe Arg Val
165 170 175
Asp Glu Val Asn Trp Ser His Trp Asn Gln Asn Leu Gly Ile Ile Asn
180 185 190
Glu Asp Pro Gly Lys Asn Glu Thr Tyr Gln Tyr Tyr Gly Phe Ser His
195 200 205
Thr Val Gly Arg Leu Arg Arg Asp Arg Trp Ser Ser Val Val Pro Arg
210 215 220
Val Val Glu Leu Asn Lys Asn Ser Asn Pro Asp Glu Val Val Val Pro
225 230 235 240
Leu Asp Ser Met Gly Asn Pro Arg Cys Asp Gly His Gln Gln Gly Tyr
245 250 255
Pro Arg Lys Trp Arg Thr Asp Asp Ala Pro Leu *
260 265 267

<210> 1048
<211> 59
<212> PRT
<213> Homo sapiens

<400> 1048
Met Trp Ser His Phe Trp Lys Val Ser Thr Gln Gly Leu Phe Val Ala
1 5 10 15
Met Phe Trp Pro Leu Ile Pro Gln Phe Val Cys Asn Cys Leu Phe Tyr
20 25 30
Trp Ala Leu Tyr Phe Asn Pro Ile Ile Asn Ile Asp Leu Val Val Lys
35 40 45
Glu Leu Arg Arg Leu Glu Thr Gln Val Leu *
50 55 58

<210> 1049
<211> 77
<212> PRT
<213> Homo sapiens

<400> 1049
Met Arg Cys Arg Cys Cys Leu Cys Ser Ser Cys Phe Trp Gly Leu Trp
1 5 10 15
Asp Pro Cys Pro Lys Ser Val Trp Ser Pro Trp Ser Ser Ser Ser Leu
20 25 30
Gly Ala Phe Ser Val Gly Ser Glu Leu Ala Ser Ala Ala Ser Ser Leu
35 40 45
Ser Pro Pro Ser Cys Ser Pro Arg Thr Ala Pro Arg Ser Thr Ala Lys
50 55 60
Leu Cys Leu Arg Trp Ser Arg Pro Gly Asn Cys Gly *
65 70 75 76

<210> 1050
<211> 474
<212> PRT
<213> Homo sapiens

<400> 1050
Met Arg Ala Leu Val Leu Leu Gly Cys Leu Leu Ala Ser Leu Leu Phe
1 5 10 15
Ser Gly Gln Ala Glu Glu Thr Glu Asp Ala Asn Glu Glu Ala Pro Leu
20 25 30
Arg Asp Arg Ser His Ile Glu Lys Thr Leu Met Leu Asn Glu Asp Lys
35 40 45
Pro Ser Asp Asp Tyr Ser Ala Val Leu Gln Arg Leu Arg Lys Ile Tyr
50 55 60
His Ser Ser Ile Lys Pro Leu Glu Gln Ser Tyr Lys Tyr Asn Glu Leu
65 70 75 80
Arg Gln His Glu Ile Thr Asp Gly Glu Ile Thr Ser Lys Pro Met Val
85 90 95
Leu Phe Leu Gly Pro Trp Ser Val Gly Lys Ser Thr Met Ile Asn Tyr
100 105 110
Leu Leu Gly Leu Glu Asn Thr Arg Tyr Gln Leu Tyr Thr Gly Ala Glu
115 120 125
Pro Thr Thr Ser Glu Phe Thr Val Leu Met His Gly Pro Lys Leu Lys
130 135 140
Thr Ile Glu Gly Ile Val Met Ala Ala Asp Ser Ala Arg Ser Phe Ser
145 150 155 160
Pro Leu Glu Lys Phe Gly Gln Asn Phe Leu Glu Lys Leu Ile Gly Ile
165 170 175
Glu Val Pro His Lys Leu Leu Glu Arg Val Thr Phe Val Asp Thr Pro
180 185 190
Gly Ile Ile Glu Asn Arg Lys Gln Gln Glu Arg Gly Tyr Pro Phe Asn
195 200 205
Asp Val Cys Gln Trp Phe Ile Asp Arg Ala Asp Leu Ile Phe Val Val
210 215 220
Phe Asp Pro Thr Lys Leu Asp Val Gly Leu Glu Leu Glu Met Leu Phe
225 230 235 240
Arg Gln Leu Lys Gly Arg Glu Ser Gln Ile Arg Ile Ile Leu Asn Lys
245 250 255
Ala Asp Asn Leu Ala Thr Gln Met Leu Met Arg Val Tyr Gly Ala Leu
260 265 270
Phe Trp Ser Leu Ala Pro Leu Ile Asn Val Thr Glu Pro Pro Arg Val
275 280 285
Tyr Val Ser Ser Phe Trp Pro Gln Glu Tyr Lys Pro Asp Thr His Gln
290 295 300

```
Glu Leu Phe Leu Gln Glu Glu Ile Ser Leu Leu Glu Asp Leu Asn Gln
305                 310                 315                 320
Val Ile Glu Asn Arg Leu Glu Asn Lys Ile Ala Phe Ile Arg Gln His
                325                 330                 335
Ala Ile Arg Val Arg Ile His Ala Leu Leu Val Asp Arg Tyr Leu Gln
            340                 345                 350
Thr Tyr Lys Asp Lys Met Thr Phe Phe Ser Asp Gly Glu Leu Val Phe
        355                 360                 365
Lys Asp Ile Val Glu Asp Pro Asp Lys Phe Tyr Ile Phe Lys Thr Ile
    370                 375                 380
Leu Ala Lys Thr Asn Val Ser Lys Phe Asp Leu Pro Asn Arg Glu Ala
385                 390                 395                 400
Tyr Lys Asp Phe Phe Gly Ile Asn Pro Ile Ser Ser Phe Lys Leu Leu
                405                 410                 415
Ser Gln Gln Cys Ser Tyr Met Gly Gly Cys Phe Leu Glu Lys Ile Glu
            420                 425                 430
Arg Ala Ile Thr Gln Glu Leu Pro Gly Leu Leu Gly Ser Leu Gly Leu
        435                 440                 445
Gly Lys Asn Pro Gly Ala Leu Asn Cys Asp Lys Thr Gly Cys Ser Glu
    450                 455                 460
Thr Pro Lys Asn Arg Tyr Arg Lys His  *
465                 470         473
```

```
<210> 1051
<211> 47
<212> PRT
<213> Homo sapiens
```

```
<400> 1051
Met Gln Arg Pro Ser Ala Trp Trp Ile Leu Phe Cys Ser Leu Asn Leu
  1               5                  10                  15
Leu Ala Arg Phe Ile Gln Cys Leu Gln Ile Val Asn Lys Glu Val His
             20                  25                  30
Phe Phe Arg Tyr Ile Lys Tyr Tyr Lys Phe Trp Glu Gly Arg  *
         35                  40                  45  46
```

```
<210> 1052
<211> 233
<212> PRT
<213> Homo sapiens
```

```
<400> 1052
Met Ala Trp Thr Pro Leu Trp Leu Thr Leu Leu Thr Leu Cys Ile Gly
  1               5                  10                  15
Ser Val Val Ser Ser Glu Leu Thr Gln Asp Pro Thr Val Ser Val Ala
             20                  25                  30
Leu Gly Gln Thr Leu Arg Ile Lys Cys Gln Gly Asp Thr Ile Arg Ser
         35                  40                  45
Tyr Tyr Ala Ser Trp Tyr Gln Gln Lys Pro Gly Gln Ala Pro Ile Leu
     50                  55                  60
Val Ile Tyr Gly Gln Asn Asn Arg Pro Ser Gly Ile Pro Gly Arg Phe
 65                  70                  75                  80
Ser Gly Ser Ser Ser Gly Asn Thr Ala Ser Leu Thr Ile Ser Gly Leu
```

```
                85                  90                  95
Gln Ala Glu Asp Glu Ala Asp Tyr Tyr Cys Cys Ser Tyr Ala Gly Arg
            100                 105                 110
Thr Thr Trp Val Phe Gly Gly Gly Thr Lys Leu Thr Val Leu Gly Gln
        115                 120                 125
Pro Lys Ala Ala Pro Ser Val Thr Leu Phe Pro Pro Ser Ser Glu Glu
    130                 135                 140
Leu Gln Ala Asn Lys Ala Thr Leu Val Cys Leu Ile Ser Asp Phe Tyr
145                 150                 155                 160
Pro Gly Ala Val Thr Val Ala Trp Lys Ala Asp Ser Ser Pro Val Lys
                165                 170                 175
Ala Gly Val Glu Thr Thr Thr Pro Ser Lys Gln Ser Asn Asn Lys Tyr
            180                 185                 190
Ala Ala Ser Ser Tyr Leu Ser Leu Thr Pro Glu Gln Trp Lys Ser His
        195                 200                 205
Arg Ser Tyr Ser Cys Gln Val Thr His Glu Gly Ser Thr Val Glu Lys
    210                 215                 220
Thr Val Ala Pro Thr Glu Cys Ser  *
225                 230     232
```

<210> 1053
<211> 147
<212> PRT
<213> Homo sapiens

<400> 1053

```
Met Gly Ala Asp Arg Gly Pro His Val Val Leu Trp Thr Leu Ile Cys
 1               5                  10                  15
Leu Pro Val Val Phe Ile Leu Ser Phe Val Val Ser Phe Tyr Tyr Gly
            20                  25                  30
Thr Ile Thr Trp Tyr Asn Ile Phe Leu Val Tyr Asn Glu Glu Arg Thr
        35                  40                  45
Phe Trp His Lys Ile Ser Tyr Cys Pro Cys Leu Val Leu Phe Tyr Pro
    50                  55                  60
Val Leu Ile Met Ala Met Ala Ser Ser Leu Gly Leu Tyr Ala Ala Val
 65                 70                  75                  80
Val Gln Leu Ser Trp Ser Trp Glu Ala Trp Trp Gln Ala Ala Arg Asp
                85                  90                  95
Met Glu Lys Gly Phe Cys Gly Trp Leu Cys Ser Lys Leu Gly Leu Glu
            100                 105                 110
Asp Cys Ser Pro Tyr Ser Ile Val Glu Leu Leu Glu Ser Asp Asn Ile
        115                 120                 125
Ser Ser Thr Leu Ser Asn Lys Asp Pro Ile Gln Glu Val Glu Thr Ser
    130                 135                 140
Thr Val  *
145 146
```

<210> 1054
<211> 123
<212> PRT
<213> Homo sapiens

<400> 1054

```
Met Tyr Val Thr Leu Val Phe Arg Val Lys Gly Ser Arg Leu Val Lys
 1               5                  10                  15
Pro Ser Leu Cys Leu Ala Leu Leu Cys Pro Ala Phe Leu Val Gly Val
            20                  25                  30
Val Arg Val Ala Glu Tyr Arg Asn His Trp Ser Asp Val Leu Ala Gly
        35                  40                  45
Phe Leu Thr Gly Ala Ala Ile Ala Thr Phe Leu Val Thr Cys Val Val
    50                  55                  60
His Asn Phe Gln Ser Arg Pro Pro Ser Gly Arg Arg Leu Ser Pro Trp
 65                 70                  75                  80
Glu Asp Leu Gly Gln Ala Pro Thr Met Asp Ser Pro Leu Glu Lys Asn
                85                  90                  95
Pro Arg Ser Ala Gly Arg Ile Arg His Arg His Gly Ser Pro His Pro
            100                 105                 110
Ser Arg Arg Thr Ala Pro Ala Val Ala Thr  *
        115                 120         122


      <210> 1055
      <211> 122
      <212> PRT
      <213> Homo sapiens


      <400> 1055
Met Leu Thr Cys Leu Phe Ser Phe Gln Gly Cys Trp Arg Ala Arg Gly
 1               5                  10                  15
Trp Gln Arg Leu Cys Glu Gly Arg Arg Gly Trp Pro Gly Val Gly Gln
            20                  25                  30
Arg Thr Leu Lys Val Ser Glu Pro Ala Pro Leu Arg Val Gly Arg Ala
        35                  40                  45
Leu Pro Gln Ala Leu Leu Gly Ala Arg Pro His Cys Val Phe Pro Gly
    50                  55                  60
Gly Glu Val Leu Gly Val Glu Ala Ala Phe Gly Ser Ser Phe Ile Leu
 65                 70                  75                  80
Ser Thr Phe Phe Leu His Gln Pro Leu Phe Phe Pro Gly Pro Lys Leu
                85                  90                  95
Arg Ala Thr Gln Tyr Leu Ile Ser Ser Asp Pro Thr His Leu Pro Ala
            100                 105                 110
Gly Arg Gly Pro Asn Ser Val Ser Met  *
        115                 120 121


      <210> 1056
      <211> 51
      <212> PRT
      <213> Homo sapiens


      <400> 1056
Met Pro Thr Lys Leu Ser Ala Val Gly Ile Leu Val Gly Thr Leu Val
 1               5                  10                  15
Ala Ile Gly Ile Phe Leu Ile Leu Ile Phe Thr His Trp Thr Met Ser
            20                  25                  30
Arg Lys Lys Asp Pro Asp Gln Pro Ala Asp Ser Val Pro Leu Lys Ala
        35                  40                  45
Thr Val  *
```

50

<210> 1057
<211> 260
<212> PRT
<213> Homo sapiens

<400> 1057
Met Glu Ala Pro Ala Gln Leu Leu Phe Leu Leu Leu Leu Trp Leu Pro
1 5 10 15
Asp Thr Thr Gly Glu Ile Val Leu Thr Gln Ser Pro Ala Thr Leu Ser
20 25 30
Leu Ser Pro Gly Glu Arg Ala Thr Leu Ser Cys Arg Ala Ser Gln Ser
35 40 45
Val Gly Ser Tyr Leu Ala Trp Tyr Gln Gln Lys Pro Gly Gln Ala Pro
50 55 60
Arg Pro Leu Ile Tyr Asp Ala Ser Asn Arg Ala Thr Gly Ile Pro Ala
65 70 75 80
Arg Phe Ser Gly Ser Gly Ser Gly Thr Asp Phe Thr Leu Thr Ile Ser
85 90 95
Ser Leu Glu Pro Glu Asp Phe Ala Val Tyr Tyr Cys Gln His Arg Asp
100 105 110
Asn Trp Pro Pro Gly Ala Thr Phe Gly Gly Gly Thr Lys Val Glu Ile
115 120 125
Lys His Thr Thr Gly Glu Ile Val Leu Thr Gln Ala Pro Gly Thr Leu
130 135 140
Ser Leu Ser Pro Gly Glu Arg Ala Thr Leu Ser Cys Arg Ala Ser Gln
145 150 155 160
Thr Ile Gly Ser Thr Tyr Leu Ala Trp Tyr Gln Gln Lys Pro Gly Lys
165 170 175
Ala Pro Lys Leu Leu Ile Tyr Trp Phe Ile Gln Phe Ala Lys Arg Gly
180 185 190
Pro Ile Lys Val Gln Cys His Arg Val Arg Gly Gln Thr Ser Leu Ser
195 200 205
Pro Ser Ala Asp Trp Ser Leu Lys Ile Leu Gln Cys Ile Ser Val Thr
210 215 220
Asn Met Gly Ala His Pro Thr Leu Leu Ala Glu Gly Pro Arg Trp Arg
225 230 235 240
Ser Asn Glu Leu Trp Leu His His Leu Ser Ser Ser Ser Arg His Leu
245 250 255
Met Ser Ser *
259

<210> 1058
<211> 52
<212> PRT
<213> Homo sapiens

<400> 1058
Met Lys Gly Leu Phe Cys Leu Trp Pro Leu Val Arg Ser Val Ser Ser
1 5 10 15
Leu Met Thr Ser Ser Thr Ser Cys Pro Ser Pro Pro Thr Leu Pro Pro
20 25 30

Trp Arg Pro Cys Leu Pro Arg Leu Arg Met Arg Val Leu Val Leu Leu
35 40 45
Ile Trp Ser *
50 51

<210> 1059
<211> 97
<212> PRT
<213> Homo sapiens

<400> 1059
Met Gly Arg Gly Ser Glu Leu Pro Val Cys Leu Ala Phe Leu Val Cys
1 5 10 15
Leu Met Ala Ala Leu Gly Cys Cys Glu Val Leu Ser Thr Val His Pro
20 25 30
Glu Glu Thr Val Leu Arg Ala Pro Pro Thr Asn Phe Gln Arg Cys Gln
35 40 45
Leu Gln Gln Gly Ser Ala Leu Val Arg Glu Thr Ala Trp Gly Val Gly
50 55 60
Arg Gly Arg Pro Ser Glu Arg Trp His Gly Glu Leu Ala Gly Gly Gly
65 70 75 80
Ser Arg Arg Asp Gly Met Glu Gly Leu Gly Pro Val Leu Leu Gly Ala
85 90 95 96
*

<210> 1060
<211> 99
<212> PRT
<213> Homo sapiens

<400> 1060
Met Asn Lys His Phe Leu Phe Leu Phe Leu Leu Tyr Cys Leu Ile Ala
1 5 10 15
Ala Val Thr Ser Leu Gln Cys Ile Thr Cys His Leu Arg Thr Arg Thr
20 25 30
Asp Arg Cys Arg Arg Gly Phe Gly Val Cys Thr Ala Gln Lys Gly Glu
35 40 45
Ala Cys Met Leu Leu Arg Ile Tyr Gln Arg Asn Thr Leu Gln Ile Ser
50 55 60
Tyr Met Val Cys Gln Lys Phe Cys Arg Asp Met Thr Phe Asp Leu Arg
65 70 75 80
Asn Arg Thr Tyr Val His Thr Cys Cys Asn Tyr Asn Tyr Cys Asn Phe
85 90 95
Lys Leu *
98

<210> 1061
<211> 64
<212> PRT
<213> Homo sapiens

<400> 1061
Met Asn Val Val Ser Leu Val Ile Leu Phe Trp Ala Ile Tyr Cys Val
1 5 10 15
Thr Ile Cys Met Asp Leu Tyr Leu Lys His Phe Cys Lys Lys Phe Phe
20 25 30
Lys Val Phe Phe Lys Cys Val Ile Ile Cys Ala Phe Lys Ser Ile Leu
35 40 45
His Phe Ser Leu Ile Cys Thr Phe Lys Lys Ile Phe Phe Phe Phe *
50 55 60 63

<210> 1062
<211> 149
<212> PRT
<213> Homo sapiens

<400> 1062
Met Tyr Leu Ser Asn Thr Thr Val Thr Ile Leu Ala Asn Leu Val Pro
1 5 10 15
Phe Thr Leu Thr Leu Ile Ser Phe Leu Leu Leu Ile Cys Ser Leu Cys
20 25 30
Lys His Leu Lys Lys Met Gln Leu His Gly Lys Gly Ser Gln Asp Pro
35 40 45
Ser Met Lys Val His Ile Lys Ala Leu Gln Thr Val Thr Ser Phe Leu
50 55 60
Leu Leu Cys Ala Ile Tyr Phe Leu Ser Met Ile Ile Ser Val Cys Asn
65 70 75 80
Phe Gly Arg Leu Glu Lys Gln Pro Val Phe Met Phe Cys Gln Ala Ile
85 90 95
Ile Phe Ser Tyr Pro Ser Thr His Pro Phe Ile Leu Ile Leu Gly Asn
100 105 110
Lys Lys Leu Lys Gln Ile Phe Leu Ser Val Leu Arg His Val Arg Tyr
115 120 125
Trp Val Lys Asp Arg Ser Leu Arg Leu His Arg Phe Thr Arg Gly Ala
130 135 140
Leu Cys Val Phe *
145 148

<210> 1063
<211> 63
<212> PRT
<213> Homo sapiens

<400> 1063
Met His Gln Leu Phe Gly Leu Phe Val Thr Leu Met Phe Ala Ser Val
1 5 10 15
Gly Gly Gly Leu Gly Gly Ile Ile Leu Val Leu Cys Leu Leu Asp Pro
20 25 30
Cys Ala Leu Trp His Trp Val Ala Pro Ser Ser Met Val Gly Gly Arg
35 40 45
Glu Ala Ser Gln Ile Leu Pro Tyr His His Gln Gly Ser Cys *
50 55 60 62

<210> 1064
<211> 92
<212> PRT
<213> Homo sapiens

<400> 1064
Met Met Leu Met Ser Leu Gly Gly Leu Leu Gly Pro Pro Leu Ser Gly
1 5 10 15
Phe Leu Arg Asp Glu Thr Gly Asp Phe Thr Ala Ser Phe Leu Leu Ser
20 25 30
Gly Ser Leu Ile Leu Ser Gly Ser Phe Ile Tyr Ile Gly Leu Pro Arg
35 40 45
Ala Leu Pro Ser Cys Gly Pro Ala Ser Pro Pro Ala Thr Pro Pro Pro
50 55 60
Glu Thr Gly Glu Leu Leu Pro Ala Pro Gln Ala Val Leu Leu Ser Pro
65 70 75 80
Gly Gly Pro Gly Ser Thr Leu Asp Thr Thr Cys *
85 90 91

<210> 1065
<211> 67
<212> PRT
<213> Homo sapiens

<400> 1065
Met Phe Leu Glu His Ala Ile His Cys Ser Leu Leu Phe Leu Ser Gln
1 5 10 15
Leu Pro Leu Leu Pro Pro Leu Val Phe Leu Leu Leu Ser His Leu Leu
20 25 30
Ser Glu Val Pro Leu Ile Gln Gln Pro Pro Ser Leu Ser Pro Tyr Pro
35 40 45
Asp Leu Leu Ser Pro Phe Ser Val Thr Arg Leu Pro Ser Asn Ile Leu
50 55 60
Cys Asn *
65 66

<210> 1066
<211> 78
<212> PRT
<213> Homo sapiens

<400> 1066
Met Gly Gln Val Pro Cys Cys Trp Ala Trp Trp Ser Leu Leu Gln Gly
1 5 10 15
Arg Gly Ser Trp Cys Glu His Lys Glu Leu Arg Gly Trp Arg Arg Pro
20 25 30
Gly Pro Gly Ala Cys Arg Arg Thr Pro Ala Arg Gly Gln Ala Gly Pro
35 40 45
Gly Ala Cys Arg Arg Thr Pro Ala Arg Gly Gln Ala Gly Pro Asp Ser

```
    50                  55                  60
Leu Ala Gly Trp Asp Leu Thr Gly Ala Pro Gly Ser Leu Gly
 65                  70                  75          78
```

<210> 1067
<211> 55
<212> PRT
<213> Homo sapiens

<400> 1067

```
Met Tyr Phe Gly Ala Tyr Ala Phe Thr Val Ala Pro Arg Leu Ala Ile
  1               5                  10                  15
Leu Gln Val Val Asn Val Ile Ser Tyr Lys Asp Ile Arg His Phe Tyr
             20                  25                  30
Leu Arg His Trp Arg Asn Glu Arg Asn Cys Ile Cys His Val Asp Gly
         35                  40                  45
Ala Leu Ile Lys Glu Gln  *
     50              54
```

<210> 1068
<211> 48
<212> PRT
<213> Homo sapiens

<400> 1068

```
Met His Val Cys Met Pro Leu Cys Leu Phe Leu Leu Ser Phe Ser Val
  1               5                  10                  15
Ser Pro Asp Pro Arg Leu Leu Arg Met Glu Arg Leu Phe Arg Gly Cys
             20                  25                  30
Ala Gln Asp Cys Pro Phe Leu Ala Leu His Gln Gly Glu Leu Trp  *
         35                  40                  45      47
```

<210> 1069
<211> 64
<212> PRT
<213> Homo sapiens

<400> 1069

```
Met Ser Asn Leu Gln Phe Ile Phe Lys Asp Phe Gly Ile Leu Ile Lys
  1               5                  10                  15
Phe Trp Tyr Leu His Ile Lys Phe Gly Phe Tyr Ile Thr Ser Cys Leu
             20                  25                  30
Leu Cys Phe Pro Pro Ser Phe Met Leu Phe Phe Gly Phe Trp Pro His
         35                  40                  45
Asp Tyr Asn Leu Arg Phe Cys Ile His Ile Thr Phe Cys His Phe  *
     50                  55                  60          63
```

<210> 1070

<211> 73
<212> PRT
<213> Homo sapiens

<400> 1070
Met Pro Ser Ile Arg Leu Gly Leu Ser His Leu Phe Leu Thr Ala Gly
1               5                   10                  15
Ile Tyr Cys Leu Leu Leu Cys Ala Arg Cys Cys Ala Leu Gly Arg Gly
            20                  25                  30
Thr Ala Trp Ala Ala Cys Pro Gly Gly Ala Cys Gly Leu Met Gly Glu
        35                  40                  45
Ala Asp Pro Ser Pro Pro His Cys Gln Gln Gly Gln Gly Lys Ser Thr
    50                  55                  60
His Arg Gly Leu Ile Pro Tyr Val  *
65                  70      72

<210> 1071
<211> 152
<212> PRT
<213> Homo sapiens

<400> 1071
Met Phe Trp Thr Met Ile Ile Leu Leu Gln Val Leu Ile Pro Ile Ser
1               5                   10                  15
Leu Tyr Val Ser Ile Glu Ile Val Lys Leu Gly Gln Ile Tyr Phe Ile
            20                  25                  30
Gln Ser Asp Val Asp Phe Tyr Asn Glu Lys Met Asp Ser Ile Val Gln
        35                  40                  45
Cys Arg Ala Leu Asn Ile Ala Glu Asp Leu Gly Gln Ile Gln Tyr Leu
    50                  55                  60
Phe Ser Asp Lys Thr Gly Thr Leu Thr Glu Asn Lys Met Val Phe Arg
65                  70                  75                  80
Arg Trp Ser Gly Gly Arg Phe Asp Tyr Cys Pro Gly Glu Lys Ala Arg
                85                  90                  95
Arg Val Glu Ser Phe Gln Glu Ala Ala Phe Glu Glu Glu His Phe Leu
            100                 105                 110
Thr Thr Gly Arg Gly Phe Leu Thr His Met Ala Asn Pro Arg Ala Pro
        115                 120                 125
Pro Leu Ala Asp Thr Phe Lys Met Gly Ala Ser Gly Arg Leu Ser Pro
    130                 135                 140
Pro Ser Leu Thr Ala Arg Gly Ala
145                 150     152

<210> 1072
<211> 113
<212> PRT
<213> Homo sapiens

<400> 1072
Met Thr Ala Gly Val Leu Trp Gly Leu Phe Gly Val Leu Gly Phe Thr
1               5                   10                  15
Gly Val Ala Leu Leu Leu Tyr Ala Leu Phe His Lys Ile Ser Gly Glu

```
            20                  25                  30
Ser Ser Ala Thr Asn Glu Pro Arg Gly Ala Ser Arg Pro Asn Pro Gln
        35                  40                  45
Glu Phe Thr Tyr Ser Ser Pro Thr Pro Asp Met Glu Glu Leu Gln Pro
    50                  55                  60
Val Tyr Val Asn Val Gly Ser Val Asp Val Asp Val Val Tyr Ser Gln
65                  70                  75                  80
Val Trp Ser Met Gln Gln Pro Glu Ser Ser Ala Asn Ile Arg Thr Leu
                85                  90                  95
Leu Glu Asn Lys Asp Ser Gln Val Ile Tyr Ser Ser Val Lys Lys Ser
            100                 105                 110     112
*
```

<210> 1073
<211> 52
<212> PRT
<213> Homo sapiens

<400> 1073

```
Met Thr Leu Cys Cys Pro Trp Ala Thr Met His Pro Ser Thr Val Leu
1               5                   10                  15
Arg Met Val Trp Ser Leu Arg Ser Arg Ala Arg Arg Trp Gly Ser Val
            20                  25                  30
Arg Thr Gly Leu Ser Trp Ser Ser Ser Ser Asp Ser Arg Ile Thr Ser
        35                  40                  45
Leu Ser Leu  *
    50  51
```

<210> 1074
<211> 78
<212> PRT
<213> Homo sapiens

<400> 1074

```
Met Phe Ser Arg Leu Tyr Ala Val Cys Met Leu Tyr Met Trp Gly Phe
1               5                   10                  15
Val Asp Lys Met Cys Val Trp Ser Val Met Gln Val Cys Tyr Cys Leu
            20                  25                  30
Val Phe Val Tyr Val Phe Leu Cys Met Val Cys Arg Val Arg Ala His
        35                  40                  45
Asp His Ile Gln Ile Leu Asp Pro Tyr Ser Arg Leu Val Leu Ser Arg
    50                  55                  60
Leu Pro Arg Leu Glu Thr Gly Lys Asp Ser Ser Ser Leu  *
65                  70                  75      77
```

<210> 1075
<211> 253
<212> PRT
<213> Homo sapiens

<400> 1075
Met Ser Ser Ser Pro Gly Leu Leu Phe Ser Ser Leu Ser His Leu Leu
1 5 10 15
Leu Asn Ser Ser Thr Leu Ala Leu Leu Thr His Arg Leu Ser Gln Met
20 25 30
Thr Cys Leu Gln Ser Leu Arg Leu Asn Arg Asn Ser Ile Gly Asp Val
35 40 45
Gly Cys Cys His Leu Ser Glu Ala Leu Arg Ala Ala Thr Ser Leu Glu
50 55 60
Glu Leu Asp Leu Ser His Asn Gln Ile Gly Asp Ala Gly Asp Gln His
65 70 75 80
Leu Ala Thr Ile Leu Pro Gly Leu Pro Glu Leu Arg Lys Ile Asp Leu
85 90 95
Ser Gly Asn Ser Ile Ser Ser Ala Gly Gly Val Gln Leu Ala Glu Ser
100 105 110
Leu Val Leu Cys Arg Arg Leu Glu Glu Leu Met Leu Gly Cys Asn Ala
115 120 125
Leu Gly Asp Pro Thr Ala Leu Gly Leu Ala Gln Glu Leu Pro Gln His
130 135 140
Leu Arg Val Leu His Leu Pro Phe Ser His Leu Gly Pro Asp Gly Ala
145 150 155 160
Leu Ser Leu Ala Gln Asp Leu Asp Gly Ser Pro His Leu Glu Glu Ile
165 170 175
Ser Leu Ala Glu Asn Asn Leu Ala Gly Gly Val Leu Arg Phe Cys Met
180 185 190
Glu Leu Pro Leu Leu Arg Gln Ile Glu Leu Ser Trp Asn Leu Leu Gly
195 200 205
Asp Glu Ala Ala Ala Glu Leu Ala Gln Val Leu Pro Gln Met Gly Arg
210 215 220
Leu Lys Arg Val Glu Tyr Glu Gly Pro Gly Glu Glu Trp Asp Gly Leu
225 230 235 240
Lys Gly Asp Leu His Pro Gly Asn Thr Lys Arg Pro Leu
245 250 253

<210> 1076
<211> 64
<212> PRT
<213> Homo sapiens

<400> 1076
Met Ser Asp Ile Ser Pro Leu Leu Tyr Glu Ile Trp Leu Gly Asp Thr
1 5 10 15
Ser Ala Gly Phe Phe Thr Phe Cys Val Thr Val Leu His Val Leu Leu
20 25 30
Leu Leu Ser Ser Val Leu His Phe Leu Cys Pro Arg Asp Thr Ser Val
35 40 45
Ile Ser Pro Phe Ile Pro Pro Leu Thr Pro Pro Gln Ser Arg Leu *
50 55 60 63

<210> 1077
<211> 147
<212> PRT
<213> Homo sapiens

<400> 1077
Met Met Lys Ser Leu Arg Val Leu Leu Val Ile Leu Trp Leu Gln Leu
1 5 10 15
Ser Trp Val Trp Ser Gln Gln Lys Glu Val Glu Gln Asn Ser Gly Pro
20 25 30
Leu Ser Val Pro Glu Gly Ala Ile Ala Ser Leu Asn Cys Thr Tyr Ser
35 40 45
Asp Arg Gly Ser Gln Ser Phe Phe Trp Tyr Arg Gln Tyr Ser Gly Lys
50 55 60
Ser Pro Glu Leu Ile Met Ser Ile Tyr Ser Asn Gly Asp Lys Glu Asp
65 70 75 80
Gly Arg Phe Thr Ala Gln Leu Asn Lys Ala Ser Gln Tyr Val Ser Leu
85 90 95
Leu Ile Arg Asp Ser Gln Pro Ser Asp Ser Ala Thr Tyr Leu Cys Ala
100 105 110
Asp Tyr Ser Gly Asn Thr Pro Leu Val Phe Gly Lys Gly Thr Arg Leu
115 120 125
Ser Val Ile Ala Asn Ile Gln Asn Pro Asp Pro Ala Leu Tyr Gln Leu
130 135 140
Arg Asp Ser
145 147

<210> 1078
<211> 55
<212> PRT
<213> Homo sapiens

<400> 1078
Met Phe Gln Gly Ser Asn Ile Leu Phe Leu Leu Pro Ser Pro Gly Ile
1 5 10 15
Thr Ser Ile Asn Asp Arg Thr Tyr Phe Leu Phe Val Met Arg Ser Asn
20 25 30
Trp Leu Phe Leu Leu Thr Cys Leu Ile Ala Phe Gln Lys Asn Asn Lys
35 40 45
Ser Leu Lys Leu Leu Lys *
50 54

<210> 1079
<211> 97
<212> PRT
<213> Homo sapiens

<400> 1079
Met Ile Pro Ala Phe Gly Ile Phe Arg Leu Leu Ile Ile Ile Leu Ile
1 5 10 15
Ile Val Leu Asp Met Gly Phe Ala Leu Tyr Arg Arg Phe Phe Val Pro
20 25 30
Glu Asp Gly Ser Pro Val Ser Phe Ala Ala His Ile Ala Gly Gly Phe
35 40 45
Ala Gly Met Ser Ile Gly Tyr Thr Val Phe Ser Cys Phe Asp Lys Ala
50 55 60

Leu Met Lys Asp Pro Arg Phe Trp Ile Ala Ile Ala Ala Tyr Leu Ala
65 70 75 80
Cys Val Leu Phe Ala Val Phe Phe Asn Ile Phe Leu Ser Pro Ala Asn
85 90 95 96
*

<210> 1080
<211> 134
<212> PRT
<213> Homo sapiens

<400> 1080
Met Leu Ser Ile Leu Leu Ala Thr Leu Thr Leu Ser Leu Lys Glu Lys
1 5 10 15
Arg Gly Glu Arg Ser Ile His Gln Pro Glu Pro Ser Glu Lys Ser Val
20 25 30
Cys Leu Pro Val Ser Gly Ala Asp Pro Phe Arg Gly Ser Arg Gly Arg
35 40 45
Gly Lys Glu Ile Arg Arg Glu Lys Asp Ile Gly Leu Leu Glu His Val
50 55 60
Gly Gln Glu Val Pro Arg Arg Ile Cys Glu Gln Leu Pro Asp Ser Lys
65 70 75 80
Ala Leu Ala Arg Pro Gln Asp Gly Pro Cys Leu Leu Asp Ile Arg Lys
85 90 95
Pro Lys Gly Gln Asn Lys Asn Thr Cys Leu Val Gly Glu Gly Ser Leu
100 105 110
Arg Gly His Gln Val Gly Gln Ile Pro Leu Val Thr His Leu Trp Arg
115 120 125
Leu Pro Gln Lys Cys *
130 133

<210> 1081
<211> 185
<212> PRT
<213> Homo sapiens

<400> 1081
Met Lys Ile Leu Val Ala Phe Leu Val Val Leu Thr Ile Phe Gly Ile
1 5 10 15
Gln Ser His Gly Tyr Glu Val Phe Asn Ile Ile Ser Pro Ser Asn Asn
20 25 30
Gly Gly Asn Val Gln Glu Thr Val Thr Ile Asp Asn Glu Lys Asn Thr
35 40 45
Ala Ile Ile Asn Ile His Ala Gly Ser Cys Ser Ser Thr Thr Ile Phe
50 55 60
Asp Tyr Lys His Gly Tyr Ile Ala Ser Arg Val Leu Ser Arg Arg Ala
65 70 75 80
Cys Phe Ile Leu Lys Met Asp His Gln Asn Ile Pro Pro Leu Asn Asn
85 90 95
Leu Gln Trp Tyr Ile Tyr Glu Lys Gln Ala Leu Asp Asn Met Phe Ser
100 105 110
Ser Lys Tyr Thr Trp Val Lys Tyr Asn Pro Leu Glu Ser Leu Ile Lys

```
        115                 120                 125
Asp Val Asp Trp Phe Leu Leu Gly Ser Pro Ile Glu Lys Leu Cys Lys
    130                 135                 140
His Ile Pro Leu Tyr Lys Gly Glu Val Val Glu Asn Thr His Asn Val
145                 150                 155                 160
Gly Ala Gly Gly Cys Ala Lys Ala Gly Leu Leu Gly Ile Leu Gly Ile
                165                 170                 175
Ser Ile Cys Ala Asp Ile His Val  *
            180             184
```

```
<210> 1082
<211> 285
<212> PRT
<213> Homo sapiens

<221> misc_feature
<222> (1)...(285)
<223> Xaa = any amino acid or nothing


<400> 1082
Met Val Ile Ala Leu Ile Ile Phe Leu Arg Ser Pro Ala Met Ala Gly
 1               5                  10                  15
Gly Leu Phe Ala Ile Glu Arg Glu Phe Phe Phe Glu Leu Gly Leu Tyr
            20                  25                  30
Asp Pro Gly Leu Gln Ile Trp Gly Gly Glu Asn Phe Glu Ile Ser Tyr
        35                  40                  45
Lys Ile Trp Gln Cys Gly Gly Lys Leu Leu Phe Xaa Pro Cys Ser Arg
    50                  55                  60
Val Gly His Ile Tyr Arg Leu Glu Gly Trp Gln Gly Asn Pro Pro Pro
 65                 70                  75                  80
Ile Tyr Val Gly Ser Ser Pro Thr Leu Lys Asn Tyr Val Arg Val Val
                85                  90                  95
Glu Val Trp Trp Asp Glu Tyr Lys Asp Tyr Phe Tyr Ala Ser Arg Pro
            100                 105                 110
Glu Ser Gln Ala Leu Pro Tyr Gly Asp Ile Ser Glu Leu Lys Lys Phe
        115                 120                 125
Arg Glu Asp His Asn Cys Lys Ser Phe Lys Trp Phe Met Glu Glu Ile
    130                 135                 140
Ala Tyr Asp Ile Thr Ser His Tyr Pro Leu Pro Pro Lys Asn Val Asp
145                 150                 155                 160
Trp Gly Glu Ile Arg Gly Phe Glu Thr Ala Tyr Cys Ile Asp Ser Met
                165                 170                 175
Gly Lys Thr Asn Gly Gly Phe Val Glu Leu Gly Pro Cys His Arg Met
            180                 185                 190
Gly Gly Asn Gln Leu Phe Arg Ile Asn Glu Ala Asn Gln Leu Met Gln
        195                 200                 205
Tyr Asp Gln Cys Leu Thr Lys Gly Ala Asp Gly Ser Lys Val Met Ile
    210                 215                 220
Thr His Cys Asn Leu Asn Glu Phe Lys Glu Trp Gln Tyr Phe Lys Asn
225                 230                 235                 240
Leu His Arg Phe Thr His Ile Pro Ser Gly Lys Cys Leu Asp Arg Ser
                245                 250                 255
Glu Val Leu His Gln Val Phe Ile Ser Asn Cys Asp Ser Ser Lys Thr
            260                 265                 270
Thr Gln Lys Trp Glu Met Asn Asn Ile His Ser Val  *
        275                 280             284
```

<210> 1083
<211> 73
<212> PRT
<213> Homo sapiens

<400> 1083
Met Phe Trp Phe Leu Asn Ile Phe Ile Leu Ile Leu Ser Lys His Ser
1 5 10 15
Ser Lys Ser Leu Ser Leu Gln Leu Pro Glu Val Leu Leu Leu Phe Leu
20 25 30
Cys Gln Phe Cys Leu Arg Leu His Pro Val Arg Gly Leu Arg Leu His
35 40 45
Phe Lys Ala Lys Leu Ala Asn His His Val Ile Cys Ile Gly Leu Gly
50 55 60
Phe Phe Leu Phe Val Ser Val Leu *
65 70 72

<210> 1084
<211> 56
<212> PRT
<213> Homo sapiens

<400> 1084
Met Ile Phe Gly Thr Asp Cys Cys Ala Leu Ser Lys Tyr Met Trp Ala
1 5 10 15
Phe Val Phe Phe Leu Ile Lys Ala Arg Trp Arg Glu Lys Asn Pro Cys
20 25 30
Phe Asp Asp Ser Leu Arg Pro Glu Gln Cys Leu Leu Asp Glu Gly Ser
35 40 45
Leu Glu Lys Arg Tyr Ser Met *
50 55

<210> 1085
<211> 68
<212> PRT
<213> Homo sapiens

<400> 1085
Met Gln Ile Phe Leu Leu Leu Tyr Ala Leu Gly Arg Phe Val Leu Leu
1 5 10 15
Val Thr Phe Ser Pro Leu Val Leu Ser Leu Ser Tyr Pro Val Leu Val
20 25 30
Ser Phe Tyr Leu Arg Tyr Pro Ser Val Leu Phe Val Phe Leu His Asn
35 40 45
Val Val Ser Leu Val Phe Gly Tyr Pro Leu Gln Asn Gln Gln Gly Leu
50 55 60
Ile His Pro *
65 67

```
<210> 1086
<211> 62
<212> PRT
<213> Homo sapiens


<400> 1086
Met Cys Pro Phe Met Pro Pro Pro Gly Leu Leu Arg Leu Phe Gln Ile
 1               5                  10                  15
Val Phe Trp Val Glu His Pro Gly Ser Val Asn Pro Phe Glu Arg Ser
            20                  25                  30
Thr Ile Ile Gly Arg Ser Ala Lys Leu Lys Lys Asp Leu Lys Ser His
        35                  40                  45
Trp Glu Pro Gly Gln Gln Ala Leu Gln Gln Gly Leu Leu  *
    50                  55                  60  61


<210> 1087
<211> 294
<212> PRT
<213> Homo sapiens


<400> 1087
Met Pro Tyr Val Thr Glu Ala Thr Arg Val Gln Leu Val Leu Pro Leu
 1               5                  10                  15
Leu Val Ala Glu Ala Ala Ala Ala Pro Ala Phe Leu Glu Ala Phe Ala
            20                  25                  30
Ala Asn Val Leu Glu Pro Arg Glu His Ala Leu Leu Thr Leu Leu Leu
        35                  40                  45
Val Tyr Gly Pro Arg Glu Gly Gly Arg Gly Ala Pro Asp Pro Phe Leu
    50                  55                  60
Gly Val Lys Ala Ala Ala Ala Glu Leu Glu Arg Arg Tyr Pro Gly Thr
65                  70                  75                  80
Arg Leu Ala Trp Leu Ala Val Arg Ala Glu Ala Pro Ser Gln Val Arg
                85                  90                  95
Leu Met Asp Val Val Ser Lys Lys His Pro Val Asp Thr Leu Phe Phe
            100                 105                 110
Leu Thr Thr Val Trp Thr Arg Pro Gly Pro Glu Val Leu Asn Arg Cys
        115                 120                 125
Arg Met Asn Ala Ile Ser Gly Trp Gln Ala Phe Phe Pro Val His Phe
    130                 135                 140
Gln Glu Phe Asn Pro Ala Leu Ser Pro Gln Arg Ser Pro Pro Gly Pro
145                 150                 155                 160
Pro Gly Ala Gly Pro Asp Pro Pro Ser Pro Pro Gly Ala Asp Pro Ser
                165                 170                 175
Arg Gly Ala Pro Ile Gly Gly Arg Phe Asp Arg Gln Ala Ser Ala Glu
            180                 185                 190
Gly Cys Phe Tyr Asn Ala Asp Tyr Leu Ala Ala Arg Ala Arg Leu Ala
        195                 200                 205
Gly Glu Leu Ala Gly Gln Glu Glu Glu Glu Ala Leu Glu Gly Leu Glu
    210                 215                 220
Val Met Asp Val Phe Leu Arg Phe Ser Gly Leu His Leu Phe Arg Ala
225                 230                 235                 240
Val Glu Pro Gly Leu Val Gln Lys Phe Ser Leu Arg Asp Cys Ser Pro
                245                 250                 255
```

Arg Leu Ser Glu Glu Leu Tyr His Arg Cys Arg Leu Ser Asn Leu Glu
260 265 270
Gly Leu Gly Gly Arg Ala Gln Leu Ala Met Ala Leu Phe Glu Gln Glu
275 280 285
Gln Ala Asn Ser Thr *
290 293

<210> 1088
<211> 477
<212> PRT
<213> Homo sapiens

<400> 1088
Met Gln Trp Lys Val Thr Leu Thr Ser Arg Trp Gly Leu Leu Arg His
1 5 10 15
Cys Gln Val Leu Ala Gly Leu Leu His Leu Gly Asn Ile Gln Phe Ala
20 25 30
Ala Ser Glu Asp Glu Ala Gln Pro Cys Gln Pro Met Asp Asp Ala Lys
35 40 45
Tyr Ser Val Arg Thr Ala Ala Ser Leu Leu Gly Leu Pro Glu Asp Val
50 55 60
Leu Leu Glu Met Val Gln Ile Lys Thr Ile Arg Ala Gly Arg Gln Gln
65 70 75 80
Gln Val Phe Arg Lys Pro Cys Ala Arg Ala Glu Cys Asp Thr Arg Arg
85 90 95
Asp Cys Leu Ala Lys Leu Ile Tyr Ala Arg Leu Phe Asp Trp Leu Val
100 105 110
Ser Val Ile Asn Ser Ser Ile Cys Ala Asp Thr Asp Ser Trp Thr Thr
115 120 125
Phe Ile Gly Leu Leu Asp Val Tyr Gly Phe Glu Ser Phe Pro Asp Asn
130 135 140
Ser Leu Glu Gln Leu Cys Ile Asn Tyr Ala Asn Glu Lys Leu Gln Gln
145 150 155 160
His Phe Val Ala His Tyr Leu Arg Ala Gln Gln Glu Glu Tyr Ala Val
165 170 175
Glu Gly Leu Glu Trp Ser Phe Ile Asn Tyr Gln Asp Asn Gln Pro Cys
180 185 190
Leu Asp Leu Ile Glu Gly Ser Pro Ile Ser Ile Cys Ser Leu Ile Asn
195 200 205
Glu Glu Cys Arg Leu Asn Arg Pro Ser Ser Ala Ala Gln Leu Gln Thr
210 215 220
Arg Ile Glu Thr Ala Leu Ala Gly Ser Pro Cys Leu Gly His Asn Lys
225 230 235 240
Leu Ser Arg Glu Pro Ser Phe Ile Val Val His Tyr Ala Gly Pro Val
245 250 255
Arg Tyr His Thr Ala Gly Leu Val Glu Lys Asn Lys Asp Pro Ile Pro
260 265 270
Pro Glu Leu Thr Arg Leu Leu Gln Gln Ser Gln Asp Pro Leu Leu Met
275 280 285
Gly Leu Phe Pro Thr Asn Pro Lys Glu Lys Thr Gln Glu Glu Pro Pro
290 295 300
Gly Gln Ser Arg Ala Pro Val Leu Thr Val Val Ser Lys Phe Lys Ala
305 310 315 320
Ser Leu Glu Gln Leu Leu Gln Val Leu His Ser Thr Thr Pro His Tyr
325 330 335
Ile Arg Cys Ile Met Pro Asn Ser Gln Gly Gln Ala Gln Thr Phe Leu

```
            340                     345                     350
Gln Glu Glu Val Leu Ser Gln Leu Glu Ala Cys Gly Leu Val Glu Thr
        355                     360                     365
Ile His Ile Ser Ala Ala Gly Phe Pro Ile Arg Val Ser His Arg Asn
    370                     375                     380
Phe Val Glu Arg Tyr Lys Leu Leu Arg Arg Leu His Pro Cys Thr Ser
385                     390                     395                 400
Ser Gly Pro Asp Ser Pro Tyr Pro Ala Lys Gly Leu Pro Glu Trp Cys
                405                     410                     415
Pro His Ser Glu Glu Ala Thr Leu Glu Pro Leu Ile Gln Asp Ile Leu
            420                     425                     430
His Thr Leu Pro Val Leu Thr Gln Ala Ala Ala Ile Thr Gly Asp Ser
        435                     440                     445
Ala Glu Ala Met Pro Ala Pro Met His Cys Gly Arg Thr Lys Val Phe
    450                     455                     460
Met Thr Asp Ser Met Leu Glu Leu Leu Glu Cys Gly Ala
465                 470                     475     477


     <210> 1089
     <211> 66
     <212> PRT
     <213> Homo sapiens


     <400> 1089
Met Ala Ala Gly Val Ser Ser Val Leu Leu Leu Leu Phe Thr Leu Met
 1               5                  10                  15
Glu Ser Gly Leu Lys His Arg Val Trp Glu Ser Trp Gln Leu Phe Thr
            20                  25                  30
Ser Trp Leu Ala Phe Cys Ser Pro Ser Phe Ser Val Val Phe Thr Cys
        35                  40                  45
Ser Tyr Ser Leu Ser Ser Trp Gly Leu Lys Gly Ile Ser Ser Arg Thr
    50                  55                  60
Arg  *
 65


     <210> 1090
     <211> 185
     <212> PRT
     <213> Homo sapiens


     <400> 1090
Met Leu Trp Leu Leu Phe Phe Leu Val Thr Ala Ile His Ala Glu Leu
 1               5                  10                  15
Cys Gln Pro Gly Ala Glu Asn Ala Phe Lys Val Arg Leu Ser Ile Arg
            20                  25                  30
Thr Ala Leu Gly Asp Lys Ala Tyr Ala Trp Asp Thr Asn Glu Glu Tyr
        35                  40                  45
Leu Phe Lys Ala Met Val Ala Phe Ser Met Arg Lys Val Pro Asn Arg
    50                  55                  60
Glu Ala Thr Glu Ile Ser His Val Leu Leu Cys Asn Val Thr Gln Arg
65                  70                  75                  80
Val Ser Phe Trp Phe Val Val Thr Asp Pro Ser Lys Asn His Thr Leu
                85                  90                  95
```

```
Pro Ala Val Glu Val Gln Ser Ala Ile Arg Met Asn Lys Asn Arg Ile
            100                 105                 110
Asn Asn Ala Phe Phe Leu Asn Asp Gln Thr Leu Glu Phe Leu Lys Ile
        115                 120                 125
Pro Ser Thr Leu Ala Pro Pro Met Asp Pro Ser Val Pro Ile Trp Ile
    130                 135                 140
Ile Ile Phe Gly Val Ile Phe Cys Ile Ile Ile Val Ala Ile Ala Leu
145                 150                 155                 160
Leu Ile Leu Ser Gly Ile Trp Gln Arg Arg Arg Lys Asn Lys Glu Pro
                165                 170                 175
Ser Glu Val Asp Asp Ala Glu Glu  *
            180                 184
```

```
     <210> 1091
     <211> 47
     <212> PRT
     <213> Homo sapiens


     <400> 1091
Met Leu Gly Gly Asn Phe Leu Met Phe Leu Pro Pro Leu Gln Arg Leu
  1               5                  10                  15
Cys Ser Asn Leu Leu Ser Tyr Val Ile Pro Asn Asp Phe Ser Val Met
             20                  25                  30
Ser Cys Phe Ile Lys Ala Ser Leu Asn Tyr Thr Leu Leu Ile  *
        35                  40                  45  46
```

```
     <210> 1092
     <211> 46
     <212> PRT
     <213> Homo sapiens


     <400> 1092
Met Val Leu Trp Asn Leu Met Leu His Ser Leu Ser Ala Val Thr Tyr
  1               5                  10                  15
Pro Pro Asp Leu Val Ser Trp Asn Leu His Phe Lys Gln Asn Pro Asp
             20                  25                  30
His Ser Pro Leu Pro Gln Leu Thr Trp Glu Val Leu Pro  *
        35                  40                  45
```

```
     <210> 1093
     <211> 64
     <212> PRT
     <213> Homo sapiens


     <400> 1093
Met Thr Val Ser Phe Cys Cys Cys Trp Ile Leu Ala Val Leu Pro Ser
  1               5                  10                  15
Pro Pro Leu Tyr Gln Asp Leu Val Gly Ser Lys Leu Glu Ile Gln Ala
             20                  25                  30
Ala Gly Asp Pro Met Pro Ala Ala Ser Arg Leu Phe His Glu Arg Gln
```

```
        35                  40                  45
Ser Leu Pro Gly Ala Pro Ala Thr Ser Ala Ser Pro Ser Val Leu  *
     50                  55                  60          63
```

<210> 1094
<211> 85
<212> PRT
<213> Homo sapiens

<400> 1094

```
Met His Phe Leu Ala Thr Phe Ala Leu Phe Phe Ile Phe Gly Val Phe
 1               5                  10                  15
Phe Leu Phe Ala Val Leu Thr Asn Leu Leu Leu Ala Glu Glu Val Asn
            20                  25                  30
Ile Arg Gly Gly Asn Phe Leu Gly Ser Phe Leu Val His Thr Leu Phe
        35                  40                  45
Leu Asp Gln Val Pro Gly Glu Ile Thr His Asp Ser His Leu Val Leu
    50                  55                  60
Ala Ile Thr Ile Asn Thr Ala Ser Pro Lys Phe Ser Ser Ser Ile Phe
65                  70                  75                  80
Phe Tyr Gln Leu  *
            84
```

<210> 1095
<211> 89
<212> PRT
<213> Homo sapiens

<400> 1095

```
Met Ala Ser His Gly Glu Glu Asp Arg His Trp Leu Arg Ala Cys Thr
 1               5                  10                  15
Trp Ile Trp Ala Leu Ser Leu Thr Leu Ser Val Ser Ser Ser Val Gly
            20                  25                  30
Trp Arg Arg Gly Gly Cys Arg Trp Leu Gly Arg Arg Asn Ala Thr Val
        35                  40                  45
Pro Arg Asn Ser Pro His Gly Thr Ser Cys Leu His Cys Val Leu Asp
    50                  55                  60
Ile Pro Ala Lys Cys Gly Arg Lys Arg Ser Gly Glu Gly Thr Phe Gln
65                  70                  75                  80
Ser Leu Leu Leu Phe Cys Thr Ala  *
                85          88
```

<210> 1096
<211> 158
<212> PRT
<213> Homo sapiens

<400> 1096

```
Met Phe Val Ile Ala Phe Leu Ser Pro Leu Ser Leu Ile Phe Leu Ala
 1               5                  10                  15
```

Lys Phe Leu Lys Lys Ala Asp Thr Arg Asp Ser Arg Gln Ala Cys Leu
20 25 30
Ala Ala Ser Leu Ala Leu Ala Leu Asn Gly Val Phe Thr Asn Thr Ile
35 40 45
Lys Leu Ile Val Gly Arg Pro Arg Pro Asp Phe Phe Tyr Arg Cys Phe
50 55 60
Pro Asp Gly Leu Ala His Ser Asp Leu Met Cys Thr Gly Asp Lys Asp
65 70 75 80
Val Val Asn Glu Gly Arg Lys Ser Phe Pro Ser Gly His Ser Ser Phe
85 90 95
Ala Phe Ala Gly Leu Ala Phe Ala Ser Phe Tyr Leu Ala Gly Lys Leu
100 105 110
His Cys Phe Thr Pro Gln Gly Arg Gly Lys Ser Trp Arg Phe Cys Ala
115 120 125
Phe Leu Ser Pro Leu Leu Phe Ala Ala Val Ile Ala Leu Ser Arg Thr
130 135 140
Cys Asp Tyr Lys His His Trp Gln Gly Pro Phe Lys Trp *
145 150 155 157

<210> 1097
<211> 88
<212> PRT
<213> Homo sapiens

<400> 1097
Met Ile Thr Thr Ser Leu Lys Ser Ser Ser Arg Leu Cys Cys Phe Arg
1 5 10 15
Arg Ser Ile Phe Phe Thr Ala Thr Cys Phe Pro Val Cys Phe Ser Val
20 25 30
Ala Met His Thr Met Pro Val Glu Pro Ser Pro Ile Leu Ile Lys Leu
35 40 45
Ala Lys Tyr Ser Leu Gly Ser Pro Gly Leu Thr Thr Ser Cys Arg Ala
50 55 60
Ala Arg Asn Cys Ser Trp Asp Thr Leu Glu Gly Cys Trp Ser Glu Glu
65 70 75 80
Glu Pro Gln Leu Gly Gly Gly *
85 87

<210> 1098
<211> 58
<212> PRT
<213> Homo sapiens

<400> 1098
Met Met Ser Gly Trp Leu Leu Arg Ala Ala Ile Cys Arg Gly Leu Leu
1 5 10 15
Ser Ser Glu Ser Leu Thr Phe Thr Ser Ala Pro His Ser Ile Ser Ile
20 25 30
Ala Val Thr Cys Arg Asp Gly Asn Leu Gln Thr Gly Tyr Arg Pro Thr
35 40 45
His Val Val Phe Leu Ser Thr Ala Arg *
50 55 57

<210> 1099
<211> 72
<212> PRT
<213> Homo sapiens

<400> 1099
Met Ala Ser Glu Pro Cys Trp Trp Ala Gly Met Leu Pro Cys Ala Cys
1 5 10 15
Ala Gly Leu Arg Arg Cys Ser His Ser Arg Phe Leu Gln Arg Gly His
20 25 30
Gly Leu His Ser Leu Met Gly Ser Leu Pro Ala Pro Ile Ser Pro Pro
35 40 45
Trp Thr His Pro Trp Gly Ile Ile Leu Pro Trp Pro Ile Arg Gly His
50 55 60
Pro Ser Val Pro Ile Arg Leu *
65 70 71

<210> 1100
<211> 47
<212> PRT
<213> Homo sapiens

<400> 1100
Met Ser Phe Phe Leu Ile Leu Gly Val Gly Ser Cys Leu Ser Tyr Ser
1 5 10 15
Leu Val Pro Leu Ile Ile Leu Ser Phe Cys His Phe Tyr Pro Glu Ser
20 25 30
Val Gly Cys Pro Asp Ala Pro Ser Pro Arg Val Arg Gly Arg Val
35 40 45 47

<210> 1101
<211> 130
<212> PRT
<213> Homo sapiens

<400> 1101
Met Arg Pro Leu Lys Pro Gly Ala Pro Leu Pro Ala Leu Phe Leu Leu
1 5 10 15
Ala Leu Ala Leu Ser Pro His Gly Ala His Gly Arg Pro Arg Gly Arg
20 25 30
Arg Gly Ala Arg Val Thr Asp Lys Glu Pro Lys Pro Leu Leu Phe Leu
35 40 45
Pro Ala Ala Gly Ala Gly Arg Thr Pro Ser Gly Ser Arg Ser Ala Glu
50 55 60
Ile Phe Pro Arg Asp Ser Asn Leu Lys Asp Lys Phe Ile Lys His Phe
65 70 75 80
Thr Gly Pro Val Thr Phe Ser Pro Glu Cys Ser Lys His Phe His Arg
85 90 95
Leu Tyr Tyr Asn Thr Arg Glu Cys Ser Thr Pro Ala Tyr Tyr Lys Arg
100 105 110

```
Cys Ala Arg Leu Leu Thr Arg Leu Ala Val Ser Pro Leu Cys Ser Gln
        115                 120                 125
Thr  *
129
```

<210> 1102
<211> 170
<212> PRT
<213> Homo sapiens

<400> 1102
```
Met Gln Phe Val Leu Leu Arg Thr Leu Ala Tyr Ile Pro Thr Pro Ile
  1               5                  10                  15
Tyr Phe Gly Ala Val Ile Asp Thr Thr Cys Met Leu Trp Gln Gln Glu
             20                  25                  30
Cys Gly Val Gln Gly Ser Cys Trp Glu Tyr Asn Val Thr Ser Phe Arg
         35                  40                  45
Phe Val Tyr Phe Gly Leu Ala Ala Val Leu Lys Tyr Val Gly Cys Ile
     50                  55                  60
Phe Ile Leu Leu Ala Trp Tyr Ser Ile Lys Asp Thr Glu Asp Glu Gln
 65                  70                  75                  80
Pro Arg Leu Arg Gln Lys Lys Ile Cys Leu Ser Thr Leu Ser Asp Thr
                 85                  90                  95
Met Thr Gln Pro Asp Ser Ala Gly Val Val Ser Cys Pro Leu Phe Thr
            100                 105                 110
Pro Asp Gly Glu Ile His Lys Lys Thr Gly Leu Arg Lys Arg Asp Pro
        115                 120                 125
Gly Gly Thr Thr Glu Pro Thr Pro Gly Pro Leu Arg Lys Arg Pro Leu
    130                 135                 140
Cys Thr Leu Glu Ala Pro Arg Leu Pro Asn Lys Ala Pro Phe Thr Leu
145                 150                 155                 160
Glu Leu Ala Leu Leu Arg Val Arg Leu  *
                165                 169
```

<210> 1103
<211> 62
<212> PRT
<213> Homo sapiens

<400> 1103
```
Met Leu Ile Ile Phe Asn Ala Val Trp Val Arg Cys Leu Lys Pro Lys
  1               5                  10                  15
Ile Pro Ala Arg Pro Thr Thr Asn Asp Thr Met Ile Ser Lys Thr Lys
             20                  25                  30
Gln His Thr Gln Tyr Thr Ser Tyr Ala Pro Ser Trp Pro Trp Leu Gly
         35                  40                  45
Pro Ala Ala Cys Gln His Gly Pro Leu Ile Ser His Thr Pro
     50                  55                  60      62
```

<210> 1104
<211> 83

<212> PRT
<213> Homo sapiens

<400> 1104
Met Lys Gln Leu Ser Pro Leu Pro Leu Pro Trp Val Leu Cys Phe Leu
1 5 10 15
Trp Lys Pro Ser Lys Leu Ser Val Leu Ser Phe Ala Ser Pro Pro Ser
20 25 30
Thr Lys Pro Ser Gln Gln Ala Gly Leu Val Cys Ser Leu Ile Arg Val
35 40 45
Ser Thr Ser Ser Thr Pro Ala Cys Thr Phe Tyr Leu Pro Val Asn Ala
50 55 60
Lys Cys Arg Ser Cys Pro Leu Asn Asn Pro Pro Trp Glu Val Pro Trp
65 70 75 80
Ile Asn *
82

<210> 1105
<211> 124
<212> PRT
<213> Homo sapiens

<400> 1105
Met Val Phe Thr Val Thr Leu Lys Leu Ala Leu Asp Thr His Tyr Trp
1 5 10 15
Thr Trp Ile Asn His Phe Val Ile Trp Gly Ser Leu Leu Phe Tyr Val
20 25 30
Val Phe Ser Leu Leu Trp Gly Gly Val Ile Trp Pro Phe Leu Asn Tyr
35 40 45
Gln Arg Met Tyr Tyr Val Phe Ile Gln Met Leu Ser Ser Gly Pro Ala
50 55 60
Trp Leu Ala Ile Val Leu Leu Val Thr Ile Ser Leu Leu Pro Asp Val
65 70 75 80
Leu Lys Lys Val Leu Cys Arg Gln Leu Trp Pro Thr Ala Thr Glu Arg
85 90 95
Val Gln Thr Lys Ser Gln Cys Leu Ser Val Glu Gln Ser Thr Ile Phe
100 105 110
Met Leu Ser Gln Thr Ser Ser Ser Leu Ser Phe *
115 120 123

<210> 1106
<211> 248
<212> PRT
<213> Homo sapiens

<400> 1106
Met Ser Phe Ser Ala Tyr Gln Thr Ala Phe Ile Cys Leu Gly Leu Leu
1 5 10 15
Val Gln Gln Ile Ile Phe Phe Leu Gly Thr Thr Ala Leu Ala Phe Leu
20 25 30
Val Leu Met Pro Val Leu His Gly Arg Asn Leu Leu Leu Phe Arg Ser
35 40 45

```
Leu Glu Ser Ser Trp Pro Phe Trp Leu Thr Leu Ala Leu Ala Val Ile
    50                  55                  60
Leu Gln Asn Met Ala Ala His Trp Val Phe Leu Glu Thr His Asp Gly
65                  70                  75                  80
His Pro Gln Leu Thr Asn Arg Arg Val Leu Tyr Ala Ala Thr Phe Leu
                85                  90                  95
Leu Phe Pro Leu Asn Val Leu Val Gly Ala Met Val Ala Thr Trp Arg
            100                 105                 110
Val Leu Leu Ser Ala Leu Tyr Asn Ala Ile His Leu Gly Gln Met Asp
        115                 120                 125
Leu Ser Leu Leu Pro Pro Arg Ala Ala Thr Leu Asp Pro Gly Tyr Tyr
    130                 135                 140
Thr Tyr Arg Asn Phe Leu Lys Ile Glu Val Ser Gln Ser His Pro Ala
145                 150                 155                 160
Met Thr Ala Phe Cys Ser Leu Leu Leu Gln Ala Gln Ser Leu Leu Pro
                165                 170                 175
Arg Thr Met Ala Ala Pro Gln Asp Ser Leu Arg Pro Gly Glu Glu Asp
            180                 185                 190
Glu Gly Met Gln Leu Leu Gln Thr Lys Asp Ser Met Ala Lys Gly Ala
        195                 200                 205
Arg Pro Gly Ala Ser Arg Gly Arg Ala Arg Trp Gly Leu Ala Tyr Thr
    210                 215                 220
Leu Leu His Asn Pro Thr Leu Gln Val Phe Arg Lys Thr Ala Leu Leu
225                 230                 235                 240
Gly Ala Asn Gly Ala Gln Pro  *
                245     247
```

<210> 1107
<211> 121
<212> PRT
<213> Homo sapiens

<400> 1107

```
Met Met Leu Ala Phe Thr Met Trp Asn Pro Trp Ile Ala Met Cys Leu
 1               5                  10                  15
Leu Gly Leu Ser Tyr Ser Leu Leu Ala Cys Ala Leu Trp Pro Met Val
            20                  25                  30
Ala Phe Val Val Pro Glu His Gln Leu Gly Thr Ala Tyr Gly Phe Met
        35                  40                  45
Gln Ser Ile Gln Asn Leu Gly Leu Ala Ile Ile Ser Ile Ile Ala Gly
    50                  55                  60
Met Ile Leu Asp Ser Arg Gly Tyr Leu Phe Leu Glu Val Phe Phe Ile
65                  70                  75                  80
Ala Cys Val Ser Leu Ser Leu Leu Ser Val Val Leu Leu Tyr Leu Val
                85                  90                  95
Asn Arg Ala Gln Gly Gly Asn Leu Asn Tyr Ser Ala Arg Gln Arg Glu
            100                 105                 110
Glu Ile Lys Phe Ser His Thr Glu  *
        115                 120
```

<210> 1108
<211> 53
<212> PRT
<213> Homo sapiens

<400> 1108
Met Phe Lys Asn Thr Ser Gly Tyr Thr Glu Arg Val Ala Val Trp Leu
1 5 10 15
Gly Val Glu Ile Phe Cys Leu Leu Met Met Ser Ser Val Leu Val Pro
20 25 30
Leu Phe Tyr Phe Leu Met Leu Phe Gly Asn Phe Leu Gln Asn Leu Ser
35 40 45
Leu Gly Ser Arg *
50 52

<210> 1109
<211> 259
<212> PRT
<213> Homo sapiens

<400> 1109
Met His Val Val Ile Val Leu Lys Ala Leu Val Ala Val Gln Ile Leu
1 5 10 15
Leu Ser Ile Lys Glu Tyr Thr Leu Glu Arg Asn His Met His Val Ile
20 25 30
Ser Val Ile Lys Val Leu Val Lys Ala Gln Thr Ser Leu Asn Ile Arg
35 40 45
Glu Tyr Thr Leu Val Lys Ser Leu Ile Ile Ala Ile Val Val Arg Lys
50 55 60
Pro Ser Val Arg Val Leu Thr Leu Phe Phe Ile Arg Glu Phe Thr Leu
65 70 75 80
Glu Lys Asn Tyr Tyr Leu Cys Thr Gln Cys Ser Lys Ser Phe Ser Gln
85 90 95
Ile Ser Asp Leu Ile Lys His Gln Arg Ile His Thr Gly Glu Lys Pro
100 105 110
Tyr Lys Cys Ser Glu Cys Arg Lys Ala Phe Ser Gln Cys Ser Ala Leu
115 120 125
Thr Leu His Gln Arg Ile His Thr Gly Lys Lys Pro Asn Pro Cys Asp
130 135 140
Glu Cys Gly Lys Ser Phe Ser Arg Arg Ser Asp Leu Ile Asn His Gln
145 150 155 160
Lys Ile His Thr Gly Glu Lys Pro Tyr Lys Cys Asp Ala Cys Gly Lys
165 170 175
Ala Phe Ser Thr Cys Thr Asp Leu Ile Glu His Gln Lys Thr His Ala
180 185 190
Glu Glu Lys Pro Tyr Gln Cys Val Gln Cys Ser Arg Ser Cys Ser Gln
195 200 205
Leu Ser Glu Leu Thr Ile His Glu Glu Val His Cys Gly Glu Asp Ser
210 215 220
Gln Asn Val Met Asn Val Arg Lys Pro Leu Val Cys Thr Pro Thr Leu
225 230 235 240
Phe Ser Thr Arg Asp Thr Val Pro Glu Lys Asn Leu Met Asn Ala Val
245 250 255
Asp Tyr *
258

<210> 1110

<211> 47
<212> PRT
<213> Homo sapiens

<400> 1110
Met Thr Cys Ser Leu Leu Ser Leu Leu Asp Ala Val Cys Ser Ser Phe
1 5 10 15
Val Gln Ala Phe Cys Ser Arg Asp Pro Glu Arg Trp Pro Ala Ile Ser
20 25 30
Pro His Ser Leu Ser Gly Ala Phe Tyr Phe Leu Asn Val Cys *
35 40 45 46

<210> 1111
<211> 93
<212> PRT
<213> Homo sapiens

<400> 1111
Met Ser Leu Arg Ala Pro Ser Val Arg Ile Phe Val Tyr Leu Leu Phe
1 5 10 15
Arg Leu His Thr Gln Arg Gly Leu Leu Ala Gly Arg Arg Gln Trp Gly
20 25 30
Pro Cys Pro Leu Ser Phe Ser His Phe Leu His Leu Ser Val Leu Ser
35 40 45
Cys Ser Thr Gln Ile Tyr Thr Glu Gly Ser Trp Pro Gly Trp Ala Ser
50 55 60
Leu Gly Ala Pro Ser Val His Trp Ala Arg Phe Pro Cys Trp Leu Gln
65 70 75 80
Ala Met Gly Ser Phe Ser Pro Leu Cys Pro Ser Cys *
85 90 92

<210> 1112
<211> 71
<212> PRT
<213> Homo sapiens

<400> 1112
Met Met Pro Thr Asn Leu Ala His Leu Val Phe Trp Gln Ala Leu Leu
1 5 10 15
Ala Ser Gly Arg Phe Ser Leu Met Glu His Tyr Pro Pro Asn Val Gln
20 25 30
Ser Asn Arg Gly Ile Thr His Tyr Met Leu Pro Arg Gly Tyr Ile Leu
35 40 45
Gly Leu Leu Tyr Ser Ser Ala Gly Asn Thr Gly Thr Ser Arg Pro Arg
50 55 60
Arg Thr His Tyr Gly Thr *
65 70

<210> 1113
<211> 47

<212> PRT
<213> Homo sapiens

<400> 1113
Met Tyr Leu Val Lys Gly Leu Leu Ile Gly Leu His Ser Ile Leu Leu
1 5 10 15
Cys Leu Arg Glu Gln Gly Gly Leu Arg Arg Val Glu Arg Asp Glu Gly
20 25 30
Thr Ala Ser Trp Tyr Ser Ser Gln Asn Thr Tyr Asn Ile Tyr *
35 40 45 46

<210> 1114
<211> 55
<212> PRT
<213> Homo sapiens

<400> 1114
Met Thr Val Leu Ser Phe Gln Tyr Glu Tyr Leu Ile Phe Leu Leu Thr
1 5 10 15
Ser Leu Thr Thr Ile Tyr Asn Thr Thr Leu Ser Arg Ser Gly Asp Gly
20 25 30
Arg Arg Thr Cys Leu Val Phe Asn Leu Arg Glu Lys Val Phe Cys Phe
35 40 45
Ser Thr Leu Gly Ile Ile *
50 54

<210> 1115
<211> 83
<212> PRT
<213> Homo sapiens

<400> 1115
Met Asn Val Ile Cys Leu Thr Leu Cys Leu Val Ser Ser Lys Cys Ser
1 5 10 15
Val Gly Gly Thr Ala Ser Phe Val Leu Leu Cys Phe Ser Leu Pro Val
20 25 30
Ser Ser Arg Arg Arg Ala Phe Gln Glu Ser Gln Gly Trp Thr Glu Pro
35 40 45
Arg Gly Gly Pro Ser Gly Leu Pro His Thr Glu Pro Gly Phe Met Ala
50 55 60
Ser Ala Ala Thr Arg Gly Leu Ser Gly Cys Gly Ser Gln Ala Ala Val
65 70 75 80
Leu Thr *
82

<210> 1116
<211> 145
<212> PRT
<213> Homo sapiens

<400> 1116
Met Val Leu Leu Val Val Gly Asn Leu Val Asn Trp Ser Phe Ala Leu
1 5 10 15
Phe Gly Leu Ile Tyr Arg Pro Arg Asp Phe Ala Ser Tyr Met Leu Gly
20 25 30
Ile Phe Ile Cys Asn Leu Leu Leu Tyr Leu Ala Phe Tyr Ile Ile Met
35 40 45
Lys Leu Arg Ser Ser Glu Lys Val Leu Pro Val Pro Leu Phe Cys Ile
50 55 60
Val Ala Thr Ala Val Met Trp Ala Ala Ala Leu Tyr Phe Phe Phe Gln
65 70 75 80
Asn Leu Ser Ser Trp Glu Gly Thr Pro Ala Glu Ser Arg Glu Lys Asn
85 90 95
Arg Glu Cys Ile Leu Leu Asp Phe Phe Asp Asp His Asp Ile Trp His
100 105 110
Phe Leu Ser Ala Thr Ala Leu Phe Phe Ser Phe Leu Asp Leu Leu Thr
115 120 125
Leu Asp Asp Asp Leu Asp Val Val Arg Arg Asp Gln Ile Pro Val Phe
130 135 140 144
*

<210> 1117
<211> 139
<212> PRT
<213> Homo sapiens

<400> 1117
Met Gly Asp Phe Ala Gly Val Asp Phe Val Phe Leu Val Val Cys Phe
1 5 10 15
Ala Gln Arg Gln Gly Ala Ala Glu Ala Val Gly Ala Val Leu Ala Val
20 25 30
Leu Leu Cys Asp Thr Leu Leu Gly Val Thr Arg Leu Glu Gly Val Ile
35 40 45
His Leu Pro Leu Tyr Phe Gly Leu Ser Gly Ile Glu Val Ile Gln Gln
50 55 60
Ala His Asn Arg Gly Ser Ser Arg Phe Gln Leu Leu Ile Arg Trp Arg
65 70 75 80
Glu Asp Glu Asp Arg Trp Cys Ser His Ser Ser Phe Asp Val His Leu
85 90 95
Gly Pro Leu Ala Glu Arg Pro His Val Ser Thr Gln Leu Leu Thr Val
100 105 110
Ile Ser Cys Lys Ile Phe Arg Leu Gln Ala Thr Asp Cys Glu Ser Lys
115 120 125
Phe Cys Pro Arg Ser Ser Ala Ala Glu Pro *
130 135 138

<210> 1118
<211> 194
<212> PRT
<213> Homo sapiens

<400> 1118
Met Cys Leu Leu Phe Leu Leu Pro Arg Phe Pro Val Ser Trp Arg Ala
1 5 10 15
Gly Val Asp Gly Ala Ala Pro Ser Ser Gln Asp Leu Trp Arg Ile Arg
20 25 30
Ser Pro Cys Gly Asp Cys Glu Gly Phe Asp Val His Ile Met Asp Asp
35 40 45
Met Ile Lys Arg Ala Leu Asp Phe Arg Glu Ser Arg Glu Ala Glu Pro
50 55 60
His Pro Leu Trp Glu Tyr Pro Cys Arg Ser Leu Ser Glu Pro Trp Gln
65 70 75 80
Ile Leu Thr Phe Asp Phe Gln Gln Pro Val Pro Leu Gln Pro Leu Cys
85 90 95
Ala Glu Gly Thr Val Glu Leu Lys Arg Pro Gly Gln Ser His Ala Ala
100 105 110
Val Leu Trp Met Glu Tyr His Leu Thr Pro Glu Cys Thr Leu Ser Thr
115 120 125
Gly Leu Leu Glu Pro Ala Asp Pro Glu Gly Gly Cys Cys Trp Asn Pro
130 135 140
His Cys Lys Gln Ala Val Tyr Phe Phe Ser Pro Ala Pro Asp Pro Arg
145 150 155 160
Ala Leu Leu Gly Gly Pro Arg Thr Val Ser Tyr Ala Val Glu Phe His
165 170 175
Pro Asp Thr Gly Asp Ile Ile Met Glu Phe Arg His Ala Asp Thr Pro
180 185 190
Asp *
193

<210> 1119
<211> 118
<212> PRT
<213> Homo sapiens

<400> 1119
Met Leu Val Leu Leu Pro Arg Ser Lys Ala Met Pro Leu Leu Ser Val
1 5 10 15
Asn Val Thr Leu Ala Phe Phe Pro Arg Asn Lys Glu Ile Val Lys Tyr
20 25 30
Leu Leu Asn Gln Gly Ala Asp Val Thr Leu Arg Ala Lys Asn Gly Tyr
35 40 45
Thr Ala Phe Asp Leu Val Met Leu Leu Asn Asp Pro Asp Ile Phe Gly
50 55 60
Gly Glu Leu Ile Gly Phe Leu Ser Val Val Thr Glu Leu Val Arg Leu
65 70 75 80
Leu Ala Ser Val Phe Met Gln Val Asn Lys Asp Ile Gly Arg Arg Ser
85 90 95
His Gln Leu Pro Leu Pro His Ser Lys Val Pro Thr Ala Leu Glu His
100 105 110
Pro Ser Ala Ala Arg *
115 117

<210> 1120
<211> 842
<212> PRT

<213> Homo sapiens

<400> 1120

```
Met Leu Trp Gly Ser Gly Lys Cys Lys Ala Leu Thr Lys Phe Lys Phe
 1               5                  10                  15
Val Phe Phe Leu Arg Leu Ser Arg Ala Gln Gly Gly Leu Phe Glu Thr
            20                  25                  30
Leu Cys Asp Gln Leu Leu Asp Ile Pro Gly Thr Ile Arg Lys Gln Thr
        35                  40                  45
Phe Met Ala Met Leu Leu Lys Leu Arg Gln Arg Val Leu Phe Leu Leu
    50                  55                  60
Asp Gly Tyr Asn Glu Phe Lys Pro Gln Asn Cys Pro Glu Ile Glu Ala
65                  70                  75                  80
Leu Ile Lys Glu Asn His Arg Phe Lys Asn Met Val Ile Val Thr Thr
                85                  90                  95
Thr Thr Glu Cys Leu Arg His Ile Arg Gln Phe Gly Ala Leu Thr Ala
            100                 105                 110
Glu Val Gly Asp Met Thr Glu Asp Ser Ala Gln Ala Leu Ile Arg Glu
        115                 120                 125
Val Leu Ile Lys Glu Leu Ala Glu Gly Leu Leu Leu Gln Ile Gln Lys
    130                 135                 140
Ser Arg Cys Leu Arg Asn Leu Met Lys Thr Pro Leu Phe Val Val Ile
145                 150                 155                 160
Thr Cys Ala Ile Gln Met Gly Glu Ser Glu Phe His Ser His Thr Gln
                165                 170                 175
Thr Thr Leu Phe His Thr Phe Tyr Asp Leu Leu Ile Gln Lys Asn Lys
            180                 185                 190
His Lys His Lys Gly Val Ala Ala Ser Asp Phe Ile Arg Ser Leu Asp
        195                 200                 205
His Cys Gly Tyr Leu Ala Leu Glu Gly Val Phe Ser His Lys Phe Asp
    210                 215                 220
Phe Glu Leu Gln Asp Val Ser Ser Val Asn Glu Asp Val Leu Leu Thr
225                 230                 235                 240
Thr Gly Leu Leu Cys Lys Tyr Thr Ala Gln Arg Phe Lys Pro Lys Tyr
                245                 250                 255
Lys Phe Phe His Lys Ser Phe Gln Glu Tyr Thr Ala Gly Arg Arg Leu
            260                 265                 270
Ser Ser Leu Leu Thr Ser His Glu Pro Glu Glu Val Thr Lys Gly Asn
        275                 280                 285
Gly Tyr Leu Gln Lys Met Val Ser Ile Ser Asp Ile Thr Ser Thr Tyr
    290                 295                 300
Ser Ser Leu Leu Arg Tyr Thr Cys Gly Ser Ser Val Glu Ala Thr Arg
305                 310                 315                 320
Ala Val Met Lys His Leu Ala Ala Val Tyr Gln His Gly Cys Leu Leu
                325                 330                 335
Gly Leu Ser Ile Ala Lys Arg Pro Leu Trp Arg Gln Glu Ser Leu Gln
            340                 345                 350
Ser Val Lys Asn Thr Thr Glu Gln Glu Ile Leu Lys Ala Ile Asn Ile
        355                 360                 365
Asn Ser Phe Val Glu Cys Gly Ile His Leu Tyr Gln Glu Ser Thr Ser
    370                 375                 380
Lys Ser Ala Leu Ser Gln Glu Phe Glu Ala Phe Phe Gln Gly Lys Ser
385                 390                 395                 400
Leu Tyr Ile Asn Ser Gly Asn Ile Pro Asp Tyr Leu Phe Asp Phe Phe
                405                 410                 415
Glu His Leu Pro Asn Cys Ala Ser Ala Leu Asp Phe Ile Lys Leu Gly
            420                 425                 430
Phe Tyr Gly Gly Ala Met Ala Ser Trp Glu Lys Ala Ala Glu Asp Thr
```

435 440 445
Gly Gly Ile His Met Glu Glu Ala Pro Glu Thr Tyr Ile Pro Ser Arg
450 455 460
Ala Val Ser Leu Phe Phe Asn Trp Lys Gln Glu Phe Arg Thr Leu Glu
465 470 475 480
Val Thr Leu Arg Asp Phe Ser Lys Leu Asn Lys Gln Asp Ile Arg Tyr
485 490 495
Leu Gly Lys Ile Phe Ser Ser Ala Thr Ser Leu Arg Leu Gln Ile Lys
500 505 510
Arg Cys Ala Gly Val Ala Gly Ser Leu Ser Leu Val Leu Ser Thr Cys
515 520 525
Lys Asn Ile Tyr Ser Leu Met Val Glu Ala Ser Pro Leu Thr Ile Glu
530 535 540
Asp Glu Arg His Ile Thr Ser Val Thr Asn Leu Lys Thr Leu Ser Ile
545 550 555 560
His Asp Leu Gln Asn Gln Arg Leu Pro Gly Gly Leu Thr Asp Ser Leu
565 570 575
Gly Asn Leu Lys Asn Leu Thr Lys Leu Ile Met Asp Asn Ile Lys Met
580 585 590
Asn Glu Glu Asp Ala Ile Lys Leu Ala Glu Gly Leu Lys Asn Leu Lys
595 600 605
Lys Met Cys Leu Phe His Leu Thr His Leu Ser Asp Ile Gly Glu Gly
610 615 620
Met Asp Tyr Ile Val Lys Ser Leu Ser Ser Glu Pro Cys Asp Leu Glu
625 630 635 640
Glu Ile Gln Leu Val Ser Cys Cys Leu Ser Ala Asn Ala Val Lys Ile
645 650 655
Leu Ala Gln Asn Leu His Asn Leu Val Lys Leu Ser Ile Leu Asp Leu
660 665 670
Ser Glu Asn Tyr Leu Glu Lys Asp Gly Asn Glu Ala Leu His Glu Leu
675 680 685
Ile Asp Arg Met Asn Val Leu Glu Gln Leu Thr Ala Leu Met Leu Pro
690 695 700
Trp Gly Cys Asp Val Gln Gly Ser Leu Ser Ser Leu Leu Lys His Leu
705 710 715 720
Glu Glu Val Pro Gln Leu Val Lys Leu Gly Leu Lys Asn Trp Arg Leu
725 730 735
Thr Asp Thr Glu Ile Arg Ile Leu Gly Ala Phe Phe Gly Lys Asn Pro
740 745 750
Leu Lys Asn Phe Gln Gln Leu Asn Leu Ala Gly Asn Arg Val Ser Ser
755 760 765
Asp Gly Trp Leu Ala Phe Met Gly Val Phe Glu Asn Leu Lys Gln Leu
770 775 780
Val Phe Phe Asp Phe Ser Thr Lys Glu Phe Leu Pro Asp Pro Ala Leu
785 790 795 800
Val Arg Lys Leu Ser Gln Val Leu Ser Lys Leu Thr Phe Leu Gln Glu
805 810 815
Ala Arg Leu Val Gly Trp Gln Phe Asp Asp Asp Asp Leu Ser Val Ile
820 825 830
Thr Gly Ala Phe Lys Leu Val Thr Ala *
835 840 841

<210> 1121
<211> 90
<212> PRT
<213> Homo sapiens

<400> 1121
Met Gly Leu Phe Phe Phe Phe Ser Gly Val Gly Ser Phe Val Gly Ser
1 5 10 15
Gly Leu Leu Ala Leu Val Ser Ile Lys Ala Ile Gly Trp Met Ser Ser
20 25 30
His Thr Asp Phe Gly Asn Ile Asn Gly Cys Tyr Leu Asn Tyr Tyr Phe
35 40 45
Phe Leu Leu Ala Ala Ile Gln Gly Ala Thr Leu Leu Leu Phe Leu Ile
50 55 60
Ile Ser Val Lys Tyr Asp His His Arg Asp His Gln Arg Ser Arg Ala
65 70 75 80
Asn Gly Val Pro Thr Ser Arg Arg Ala *
85 89

<210> 1122
<211> 129
<212> PRT
<213> Homo sapiens

<400> 1122
Met Phe Leu Leu Phe Trp Phe Ile Leu Ser Glu Gly Cys Pro Leu Leu
1 5 10 15
Glu Gln Leu Asn Ile Ser Trp Cys Asp Gln Val Thr Lys Asp Gly Ile
20 25 30
Gln Ala Leu Val Arg Gly Cys Gly Gly Leu Lys Ala Leu Phe Leu Lys
35 40 45
Gly Cys Thr Gln Leu Glu Asp Glu Ala Leu Lys Tyr Ile Gly Ala His
50 55 60
Cys Pro Glu Leu Val Thr Leu Asn Leu Gln Thr Cys Leu Gln Ile Thr
65 70 75 80
Asp Glu Gly Leu Ile Thr Ile Cys Arg Gly Cys His Lys Leu Gln Ser
85 90 95
Leu Cys Ala Ser Gly Cys Ser Asn Ile Thr Asp Ala Ile Leu Asn Ala
100 105 110
Leu Ser Gln Asn Cys Pro Arg Leu Ile Ile Leu Glu Val Ala Arg Cys
115 120 125
Ser
129

<210> 1123
<211> 243
<212> PRT
<213> Homo sapiens

<400> 1123
Met Ala Ala Ala Leu Trp Gly Phe Phe Pro Val Leu Leu Leu Leu Leu
1 5 10 15
Leu Ser Gly Asp Val Gln Ser Ser Glu Val Pro Gly Ala Ala Ala Glu
20 25 30
Gly Ser Gly Gly Ser Gly Val Gly Ile Gly Asp Arg Phe Lys Ile Glu
35 40 45
Gly Arg Ala Val Val Pro Gly Val Lys Pro Gln Asp Trp Ile Ser Ala

```
                50                  55                  60
Ala Arg Val Leu Val Asp Gly Glu Glu His Val Gly Phe Leu Lys Thr
 65                 70                  75                  80
Asp Gly Ser Phe Val Val His Asp Ile Pro Ser Gly Ser Tyr Val Val
                    85                  90                  95
Glu Val Val Ser Pro Ala Tyr Arg Phe Asp Pro Val Arg Val Asp Ile
            100                 105                 110
Thr Ser Lys Gly Lys Met Arg Ala Arg Tyr Val Asn Tyr Ile Lys Thr
        115                 120                 125
Ser Glu Val Val Arg Leu Pro Tyr Pro Leu Gln Met Lys Ser Ser Gly
    130                 135                 140
Pro Pro Ser Tyr Phe Ile Lys Arg Glu Ser Trp Gly Trp Thr Asp Phe
145                 150                 155                 160
Leu Met Asn Pro Met Val Met Met Met Val Leu Pro Leu Leu Ile Phe
                165                 170                 175
Val Leu Leu Pro Lys Val Val Asn Thr Ser Asp Pro Asp Met Arg Arg
            180                 185                 190
Glu Met Glu Gln Ser Met Asn Met Leu Asn Ser Asn His Glu Leu Pro
        195                 200                 205
Asp Val Ser Glu Phe Met Thr Arg Leu Phe Ser Ser Lys Ser Ser Gly
    210                 215                 220
Lys Ser Ser Ser Gly Ser Ser Lys Thr Gly Lys Ser Gly Ala Gly Lys
225                 230                 235                 240
Arg Arg  *
    242
```

```
<210> 1124
<211> 71
<212> PRT
<213> Homo sapiens
```

```
<400> 1124
Met Leu Ser Tyr Ala His Ile Thr Leu Ala Val Leu Arg Ile Pro Ser
  1               5                  10                  15
Ala Thr Gly Cys Trp Arg Ala Phe Phe Thr Cys Ala Ser His Leu Thr
             20                  25                  30
Val Val Thr Val Phe Tyr Thr Ala Leu Leu Phe Met Tyr Val Arg Pro
         35                  40                  45
Gln Ala Ile Asp Ser Arg Ser Ser Asn Lys Leu Ile Ser Val Leu Tyr
     50                  55                  60
Thr Val Ile Thr Pro Ser Val
 65                  70  71
```

```
<210> 1125
<211> 48
<212> PRT
<213> Homo sapiens
```

```
<400> 1125
Met Pro Thr Leu Gly Asp Ala Leu Ile Leu Tyr Leu His Leu Val Leu
  1               5                  10                  15
Gly Val Ala Gly Val Leu Gln Pro Pro Gly Pro Arg Pro Ser Gln Ala
             20                  25                  30
```

```
Leu Gly Pro Thr Gly Asp Arg Ala Pro Gly Lys Trp Asn Arg Ser  *
        35                  40                  45      47



     <210> 1126
     <211> 159
     <212> PRT
     <213> Homo sapiens


     <400> 1126
Met Phe Leu Ile Val Leu Pro Leu Glu Ser Met Ala His Gly Leu Phe
  1               5                  10                  15
His Glu Leu Gly Asn Cys Leu Gly Gly Thr Ser Val Gly Tyr Ala Ile
             20                  25                  30
Val Ile Pro Thr Asn Phe Cys Ser Pro Asp Gly Gln Pro Thr Leu Leu
        35                  40                  45
Pro Pro Glu His Val Gln Glu Leu Asn Leu Arg Ser Thr Gly Met Leu
    50                  55                  60
Asn Ala Ile Gln Arg Phe Phe Ala Tyr His Met Ile Glu Thr Tyr Gly
 65                  70                  75                  80
Cys Asp Tyr Ser Thr Ser Gly Leu Ser Phe Asp Thr Leu His Ser Lys
                 85                  90                  95
Leu Lys Ala Phe Leu Glu Leu Arg Thr Val Asp Gly Pro Arg His Asp
            100                 105                 110
Thr Tyr Ile Leu Tyr Tyr Ser Gly His Thr His Gly Thr Gly Glu Trp
        115                 120                 125
Ala Leu Ala Gly Gly Asp Thr Leu Arg Leu Asp Thr Leu Ile Glu Trp
    130                 135                 140
Trp Arg Glu Lys Asn Gly Ser Phe Cys Ser Pro Pro Tyr Tyr Arg
145                 150                 155             159



     <210> 1127
     <211> 76
     <212> PRT
     <213> Homo sapiens


     <400> 1127
Met Thr Gly Pro Arg Pro Met Ile Leu His Phe Ile Leu Val Ala Ser
  1               5                  10                  15
Ala Ser Cys Trp Glu Val Leu Phe Cys Cys Trp Gln Pro Cys Pro Leu
             20                  25                  30
Gly Ile His Ala Thr Ser Asn Ser Pro Ser Gln Leu Gln Gln Leu Ser
        35                  40                  45
Cys Thr Lys Leu Pro Leu Met Phe Arg Arg Ile Leu Glu Asp Thr Ile
    50                  55                  60
Phe Ala Ile Leu Tyr His Ile Ala Thr Ile Phe  *
 65                  70                  75



     <210> 1128
     <211> 140
     <212> PRT
     <213> Homo sapiens
```

<400> 1128
Met Gly Ala Gly Leu Ala Val Val Pro Leu Met Gly Leu Leu Glu Ser
1 5 10 15
Ile Ala Val Ala Lys Ala Phe Ala Ser Gln Asn Asn Tyr Arg Ile Asp
20 25 30
Ala Asn Gln Glu Leu Leu Ala Ile Gly Leu Thr Asn Met Leu Gly Ser
35 40 45
Leu Val Ser Ser Tyr Pro Val Thr Gly Ser Phe Gly Arg Thr Ala Val
50 55 60
Asn Ala Gln Ser Gly Val Cys Thr Pro Ala Glu Gly Leu Val Thr Glu
65 70 75 80
Val Leu Val Leu Leu Ser Leu Asp Tyr Leu Thr Ser Leu Phe Tyr Tyr
85 90 95
Ile Pro Lys Ser Ala Leu Ala Ala Val Ile Ile Met Ala Val Ala Pro
100 105 110
Leu Phe Asp Thr Lys Ile Phe Arg Thr Leu Trp Arg Val Lys Arg Leu
115 120 125
Asp Leu Leu Ser Leu Ser Val Thr Phe Leu Leu Cys
130 135 140

<210> 1129
<211> 116
<212> PRT
<213> Homo sapiens

<400> 1129
Met Ala Glu Ala Phe Pro Phe Phe Ser Pro Phe Leu Gly Trp Leu Gly
1 5 10 15
Val Phe Leu Thr Gly Ser Asp Thr Ser Ser Asn Ala Leu Phe Ser Ser
20 25 30
Leu Gln Ala Thr Thr Ala His Gln Ile Gly Val Ser Asp Val Leu Leu
35 40 45
Val Ala Ala Asn Thr Ser Gly Gly Val Thr Gly Lys Met Ile Ser Pro
50 55 60
Gln Ser Ile Ala Val Ala Cys Ala Ala Thr Gly Leu Val Gly Lys Glu
65 70 75 80
Ser Asp Leu Phe Arg Phe Thr Leu Lys His Ser Leu Phe Phe Ala Thr
85 90 95
Ile Val Gly Leu Ile Thr Leu Ala Gln Ala Tyr Trp Phe Thr Gly Met
100 105 110
Leu Val His *
115

<210> 1130
<211> 81
<212> PRT
<213> Homo sapiens

<400> 1130
Met Asn Lys Leu Leu Val Ala Ala Thr Ala Ile Leu Phe Ser Leu Gly
1 5 10 15

Cys His Glu Lys Cys Lys Ile Phe Phe Leu Lys Ser Ile Ser Ser Pro
20 25 30
Gln Ser Leu Phe Leu Ala Asp Leu Cys Ala Ser Glu Pro Tyr Leu Leu
35 40 45
Phe Leu Asn Ala Val Leu Ser Ala Cys Asn Thr Ile Ser Phe Ile Ser
50 55 60
Val Pro Glu Ser Ser Gly Phe Ala Pro Ser Pro Pro Ala Ile Leu Leu
65 70 75 80
Leu
81

<210> 1131
<211> 46
<212> PRT
<213> Homo sapiens

<400> 1131
Met Cys Cys Trp Ile Trp Phe Ala Ser Ile Leu Leu Arg Ile Phe Ala
1 5 10 15
Leu Met Phe Ile Arg Asp Ile Gly Leu Lys Phe Ser Phe Phe Val Val
20 25 30
Ser Leu Pro Gly Phe Gly Ile Arg Met Met Leu Ala Ser *
35 40 45

<210> 1132
<211> 46
<212> PRT
<213> Homo sapiens

<400> 1132
Met Ser Gln Glu Pro Gly Arg Arg His Ser Lys Leu Thr Leu Thr Ala
1 5 10 15
Ser Arg Met Ala Pro Cys Leu Trp Val Trp Thr Ser Leu Cys Gln Ala
20 25 30
Trp Ser Met Ser Met Gly Ser Leu Ser Met Gln Thr Thr *
35 40 45

<210> 1133
<211> 87
<212> PRT
<213> Homo sapiens

<400> 1133
Met His Ser His Gly Val Ser Tyr Trp Thr Val Arg Thr Val Ile Trp
1 5 10 15
Pro Ile Ser Ser Leu Val Ser Lys Ile Thr Thr Trp Glu Phe Asn Glu
20 25 30
Val Thr Ser Met Ser Glu His Leu Lys Ser Cys Pro Phe Asn Ile Val
35 40 45
Glu His Lys Ser Asp Pro Ile Leu Leu Thr Ser Met Cys His Pro Arg

```
    50                  55                  60
Glu Gln Ala Arg Glu Ser Leu Leu Ser Thr Phe Arg Ile Arg Pro Arg
 65                 70                  75                  80
Gly Arg Tyr Val Ser Tyr  *
                85  86
```

<210> 1134
<211> 57
<212> PRT
<213> Homo sapiens

<400> 1134

```
Met Glu Ala His Gln Ser Phe Lys His Lys Ser Cys Thr Trp Ala Ile
  1               5                  10                  15
Thr Val Trp Phe His Phe Val Cys Phe Leu Asn Thr Phe Ser Cys Phe
             20                  25                  30
Phe Asn Lys Leu Ser Pro Ile Leu Glu Ser Leu Val Val Gly Ser Ile
         35                  40                  45
Ser Arg His Leu Leu Arg Glu Leu  *
     50                  55  56
```

<210> 1135
<211> 57
<212> PRT
<213> Homo sapiens

<400> 1135

```
Met Glu Ala His Gln Ser Phe Lys His Lys Ser Cys Thr Trp Ala Ile
  1               5                  10                  15
Thr Val Trp Phe His Phe Val Cys Phe Leu Asn Thr Phe Ser Cys Phe
             20                  25                  30
Phe Asn Lys Leu Ser Pro Ile Leu Glu Ser Leu Val Val Gly Ser Ile
         35                  40                  45
Ser Arg His Leu Leu Arg Glu Leu  *
     50                  55  56
```

<210> 1136
<211> 105
<212> PRT
<213> Homo sapiens

<400> 1136

```
Met Pro Phe Ala Gln Thr Gly Leu Gln Leu Leu Leu Arg Leu Cys Arg
  1               5                  10                  15
Val Leu His Val Leu Arg Leu Leu Gly Met Leu Arg Glu Gln Met His
             20                  25                  30
Leu Leu Arg Glu Lys Leu Leu Asp Leu Leu Pro Pro Glu Leu Cys Gln
         35                  40                  45
Arg Val Pro Arg Ala Ala Thr Ala Lys Gly His Lys Arg Arg Ala Ala
     50                  55                  60
```

```
Ala Val Pro Asp Asp Gly Thr Asp Leu Leu Pro Gln Gly Met Arg Thr
 65                 70                  75                  80
Ala Cys Thr Thr Arg Arg Ile Phe Lys Tyr Asn Thr Glu Pro Phe Ala
                85                  90                  95
Ala Phe Leu Phe Ile Leu Asn Met  *
            100             104
```

<210> 1137
<211> 52
<212> PRT
<213> Homo sapiens

<400> 1137

```
Met Val Gly Phe Tyr Leu Gln Ser Val Leu Tyr Phe Tyr Phe Ser Gln
  1               5                  10                  15
Leu Ile Tyr Leu Gly Asp His Ala Lys Ser Val Asn Ile Val Thr Ser
            20                  25                  30
Phe Ile Leu Thr Ala Ala Tyr Val Asn Asn Ser Lys Met His His Thr
        35                  40                  45
Val Phe Asn  *
    50  51
```

<210> 1138
<211> 187
<212> PRT
<213> Homo sapiens

<400> 1138

```
Met Gln Pro Ile Val Ala Lys Ala Leu Val Val Leu Leu Glu Val His
  1               5                  10                  15
Pro Leu Gln Asp Gln Ala Glu Ser Gly Arg Leu Gly His Val His Leu
            20                  25                  30
Leu Cys Ala Pro Ala Ala Leu Gln His Ala Leu Arg Gly Ile Thr Leu
        35                  40                  45
His Asn Gly His His Gln Ala Asp His Leu Pro Asp Leu Met His His
    50                  55                  60
Glu Ala Leu Ala Leu His Pro Asp His Arg Lys Leu Gln Ala Leu Pro
 65                 70                  75                  80
His Lys Gly Phe Leu Ala Val His Leu Gln Asp Val Ala Ala Gly Thr
                85                  90                  95
Gly Ile Leu Arg Pro Leu Leu Arg Gly Glu Ile Val Glu Val Val Arg
            100                 105                 110
Ala Leu Val Ala Gly Gln Glu Pro Val Asp Leu Leu Gln Arg Leu Gly
        115                 120                 125
Ala Gln Ala Val Gly Leu Ile Leu Asn Val Pro Val Leu Val Arg Lys
    130                 135                 140
Gly Lys Arg Gly Gln Gln Val Ala Ile Gly Pro Gly Ile Thr Ser Val
145                 150                 155                 160
Leu Gly Val Lys Pro Ala Arg Asp Pro Leu Gln Ser Gln Asn Pro Asn
                165                 170                 175
Val Arg Gly Lys Val Ala Val Asp Leu Phe  *
            180                 185 186
```

<210> 1139
<211> 109
<212> PRT
<213> Homo sapiens

<400> 1139
Met Trp Gln Lys Ser Leu Leu Ile Leu Ser Phe Arg Val Ser Phe Pro
1 5 10 15
Leu Phe Leu Thr Tyr Asn Tyr Lys Leu Leu Ser Ile Arg Arg Thr Arg
20 25 30
Pro Leu Ser Ser Phe Phe Ser Lys Leu Leu Gln Ile Ala Val Asn Ser
35 40 45
Ile Asn Ser Leu Phe Ser Ala Gly Lys Val Ala Phe Ser Lys His Val
50 55 60
Cys Leu Leu Pro Gly Gly Leu Lys Ser Met Ile Tyr Cys Ser Ser Met
65 70 75 80
Cys Leu Lys Gln Leu Leu Arg Ser Phe Lys Gln Glu Ser Ser Lys Gly
85 90 95
Ser Val Leu Ile Met Val Leu Val Phe Leu Gln Ile *
100 105 108

<210> 1140
<211> 83
<212> PRT
<213> Homo sapiens

<400> 1140
Met Pro Ala Pro Thr Ala Trp Leu Leu Pro Ala Val Ser Thr Cys Ser
1 5 10 15
Asn Leu Arg Ala Lys Ala Gly Val Ile Leu Gly Thr Ile Thr Thr Arg
20 25 30
Pro Tyr Val His Thr Trp Gly Ser Ala Asp Met Ala Thr Pro Tyr His
35 40 45
Leu Gly Pro Phe Trp Thr Leu Gly Thr Asp Lys His Arg Arg Glu Ala
50 55 60
Asn Arg Gly Gln Arg Ala Ile Trp Gly Trp Pro Thr Gly Pro Pro Trp
65 70 75 80
His Leu *
82

<210> 1141
<211> 58
<212> PRT
<213> Homo sapiens

<400> 1141
Met Tyr Gln Trp Gly Ser Ser Ile Ile Leu Ile Leu Trp Pro Leu Ser
1 5 10 15
Met Asn Ile Gly Cys Tyr Ser Ile Tyr Leu Lys Met Val Met Leu Leu
20 25 30

Ser Ser Lys Phe Ser Trp Lys Ser Phe Ser Lys Leu Gln Phe Leu Leu
35 40 45
Leu Leu Lys Phe Arg Tyr Met Cys Ile *
50 55 57

<210> 1142
<211> 46
<212> PRT
<213> Homo sapiens

<400> 1142
Met Asn Pro His Leu Gly Val Phe Leu Val Leu Val Ser Phe Phe Leu
1 5 10 15
Ser Leu Leu Asp Ser Gln Leu His Ser Trp Ile Val Leu His Asn Ser
20 25 30
Pro Ser Ser Arg Met Trp Lys Ser Ile Ile Phe Phe Leu *
35 40 45

<210> 1143
<211> 58
<212> PRT
<213> Homo sapiens

<400> 1143
Met Leu Trp Ala Leu Ile Arg Ala Ala Leu Ala Gln Leu His Thr Glu
1 5 10 15
Glu Pro Lys Lys Arg Lys Glu Glu Lys Met Ser Pro Ala Leu Ser Pro
20 25 30
Pro Leu Pro Ser Val Pro Ile Ser Leu Gly Gln Asn Asn Arg Lys Arg
35 40 45
Arg Ser His Leu Ser Leu Leu Leu Gln *
50 55 57

<210> 1144
<211> 147
<212> PRT
<213> Homo sapiens

<400> 1144
Met Ala Tyr Thr Met Ile Pro Val Leu His Phe Phe Cys Cys Glu Thr
1 5 10 15
Ser Ser Leu Val Arg Thr Lys Val Val Trp Glu Ala Ile Asn Met Val
20 25 30
Phe Ala Lys Ser Met Asn Gly Gly Pro Asp Arg Cys Ile Ala Val Arg
35 40 45
Gln Val Lys Phe Leu Phe Arg Lys Val Ser Phe Ser Glu Lys Ile Asp
50 55 60
His Cys Pro Leu His Asp Gly Asn Ile Leu Leu Pro Gly Pro Trp Glu
65 70 75 80
Met Ala Pro Tyr Trp Gly Leu Asn Ile Ser Leu Cys His Leu Gln Phe

```
                85                                  90                                  95
Arg His Ser Ile Val Ser Leu Ala Arg Cys Ser Leu Gly Glu Gly Gln
            100                     105                     110
Ser Met Leu Trp Cys Pro Cys Leu Thr Ser Ile Ser Val Asp Met Ala
        115                     120                     125
Thr Leu Tyr Ile Asn Ala Ser Ser Ser Leu Ser Ser Lys Gly Lys Lys
    130                     135                     140
Ala Asp  *
145 146
```

<210> 1145
<211> 103
<212> PRT
<213> Homo sapiens

<400> 1145

```
Met Ala Trp Ile Pro Leu Phe Leu Gly Val Leu Ala Tyr Cys Thr Gly
  1               5                  10                  15
Ser Val Ala Ser Tyr Glu Leu Thr Gln Pro Pro Ser Val Ser Val Ser
             20                  25                  30
Pro Gly Lys Thr Ala Ser Ile Thr Cys Ser Gly Asp Lys Leu Gly Asp
         35                  40                  45
Lys Tyr Ala Ser Trp Tyr Gln Gln Lys Ala Gly Gln Ser Pro Val Leu
     50                  55                  60
Val Ile Tyr Glu Asp Ser Arg Arg Pro Ser Gly Ile His Lys Arg Phe
 65                  70                  75                  80
Tyr Gly Ser Asn Ser Gly Thr Thr Ala Thr Leu Thr Ile Ser Gly Thr
                 85                  90                  95
Gln Ala Met Asp Glu Gly  *
            100     102
```

<210> 1146
<211> 77
<212> PRT
<213> Homo sapiens

<400> 1146

```
Met Pro Leu Leu His Gly Val Tyr Leu Ala Arg Arg Ser Leu Ile Cys
  1               5                  10                  15
Ile Ser Phe Cys His Leu Cys Val Leu Ser Ile Gly Leu Arg Val Ile
             20                  25                  30
Val Cys Val Val Gly Ile Ser Glu Asp Arg Lys Arg Ser Ala Ser Ala
         35                  40                  45
Pro Thr Leu Gly Ile Val Pro Leu His Ala Ser Leu His Gln His Cys
     50                  55                  60
Ala Pro Asn Gln Ser Asn Pro Cys Ser Trp His Leu  *
 65                  70                  75  76
```

<210> 1147
<211> 118
<212> PRT

<213> Homo sapiens

<400> 1147

```
Met Asn Pro Ser Ala Ser Leu Val Cys Leu Leu Phe Ala Phe Ser Ser
 1               5                  10                  15
Cys Arg Ile Trp Ser Val Leu Cys Gln Leu Cys Val Pro Ser Pro Trp
            20                  25                  30
Pro Ser Pro Leu Cys Leu Cys Pro Gln Thr Asp Val Ala Pro Ile Cys
        35                  40                  45
Ala Val Gln Pro Ser Leu Phe Cys Leu Gly Ser Arg Glu Pro Leu Trp
    50                  55                  60
Thr Val Leu Val Gly Ser Cys Pro Leu Arg Ala Phe Thr Asn Leu Ser
 65                 70                  75                      80
Val Arg Pro Pro Pro Gly His His Ser Ile His Leu Leu Thr Trp Leu
                85                  90                      95
Ala Ser Ser Ser Ala Ala Ala Thr Thr Ala Ala Ser Thr Ala Ser Gly
            100                 105                 110
Ala Pro His Ser Val  *
        115     117
```

<210> 1148
<211> 399
<212> PRT
<213> Homo sapiens

<400> 1148

```
Met Trp Ala Ala Val Gly Gly Phe Leu Phe Ala Pro Arg Cys Phe Leu
 1               5                  10                  15
Leu Pro Trp Pro Leu Arg Ala Pro Leu Ser Ser Leu Phe Val Leu Pro
            20                  25                  30
Arg Leu Leu Leu Trp Pro Ile Pro Tyr Pro Val Leu Ala Ser Val Cys
        35                  40                  45
Pro Cys Val Pro Gly Gly Arg Phe Phe Gly Pro Leu Tyr Pro Arg Asp
    50                  55                  60
Leu Arg Leu Leu Arg Cys Val Pro Gly Glu Leu Thr Gly Ala Ala Pro
 65                 70                  75                      80
Arg Thr Leu Pro Gly Cys Asp Leu Asn Cys Leu Gly Leu Gly Arg Glu
                85                  90                      95
Ala Ala Val Pro Arg Leu Leu Arg Leu Thr Arg Asp Pro Ala Arg Pro
            100                 105                 110
Ser Cys Arg Thr Leu Gly Val His Ala Val Pro Arg Arg Ala Phe Gly
        115                 120                 125
Phe Tyr Ala Val Pro Arg Arg Asp Pro Arg Phe Tyr Ala Val Pro Arg
    130                 135                 140
Arg Val Pro Arg Leu Tyr Ala Val Pro His Pro Ala Leu Arg Val Tyr
145                 150                 155                     160
Ala Val Pro Arg Arg Thr Phe Arg Val Tyr Ala Val Pro His Pro Ala
                165                 170                 175
Leu Arg Val Tyr Ala Val Pro Arg Arg Ala Leu Gly Leu Tyr Val Val
            180                 185                 190
Pro Gln Arg Ala Leu Arg Val Tyr Ala Val Pro Arg Arg Thr Phe Arg
        195                 200                 205
Val Tyr Ala Val Pro His Pro Ala Leu Arg Leu Tyr Ala Val Ala Arg
    210                 215                 220
Arg Ala Leu Arg Phe Tyr Val Val Pro Gln Arg Ala Leu Arg Val Tyr
```

```
225                 230                 235                 240
Ala Val Pro Arg Leu Pro Gly Arg Ala Thr Phe Arg Asp Leu Arg Pro
                245                 250                 255
Leu Leu Arg Leu Leu Leu Pro Leu Gly Gly Arg Arg Val Leu Gly Leu
            260                 265                 270
Pro Leu Ser Leu Pro Ala Gly Leu Ala Leu Arg Ala Ala Ser Arg Ala
        275                 280                 285
Arg Pro Leu His Leu Leu Arg Ala Ala Cys Leu Leu Pro Ser Leu Gly
    290                 295                 300
His Leu Gly Thr Leu Arg Gly Ser Leu Leu Gly Leu Ser Leu Ala Val
305                 310                 315                 320
Arg Pro Pro Arg Ala Pro Arg Leu Gly Leu Arg Ala Pro Val Trp Pro
                325                 330                 335
Ala Ala Ser Cys Leu Leu His Ser Gly Gly Ala Pro Arg Arg Leu Leu
            340                 345                 350
Cys Ala Leu Ala Pro Leu Arg Pro Phe Cys Leu Pro Ala Arg Gly Ser
        355                 360                 365
Trp Leu Ser Gly Ser Leu Ser Gln Arg Arg Gly Asp Leu Arg Arg Pro
    370                 375                 380
Leu Gly Thr Arg Gly Asn Pro Leu Arg Leu Arg Gly Leu Gly His
385                 390                 395             399
```

<210> 1149
<211> 67
<212> PRT
<213> Homo sapiens

<400> 1149

```
Met Pro Ser Tyr Phe Lys Thr Cys Ser Leu Phe Thr Leu Leu Ser Ser
 1               5                  10                  15
Val Phe Leu Val Cys Ile Trp Ile Phe Lys Thr Asn Ile Lys Ser Ser
            20                  25                  30
Val Ser Glu Ser Pro Pro Asp Ser Gly Leu Gly Gln Val Thr Ala Val
        35                  40                  45
Tyr Gln Val Gln Cys Leu Cys Trp Ala Lys Asp Cys Asn Tyr Pro Ile
    50                  55                  60
Cys Ser  *
 65  66
```

<210> 1150
<211> 70
<212> PRT
<213> Homo sapiens

<400> 1150

```
Met Leu Val Ser Lys Leu Met Leu Gln Ile Val Met Ala Val Pro His
 1               5                  10                  15
Tyr Ile Met Pro Val Glu Met Lys Asn Gln Ser Leu Ile Pro Leu Leu
            20                  25                  30
Leu Glu Ala Arg Ala Asp Pro Thr Ile Lys Asn Lys His Gly Glu Ser
        35                  40                  45
Ser Leu Asp Ile Ala Arg Arg Leu Lys Phe Ser Gln Ile Glu Leu Met
    50                  55                  60
```

```
Leu Arg Lys Ala Leu  *
 65              69


     <210> 1151
     <211> 48
     <212> PRT
     <213> Homo sapiens


     <400> 1151
Met Gly Ala Gly Cys Thr Pro Val Val Leu Gly Ala Ala Leu Trp Leu
  1               5                  10                  15
Trp Arg Trp Phe Ser Arg Trp Gly Leu Gly Gly Leu Cys Trp Arg Pro
             20                  25                  30
Cys Thr Cys Thr Pro Cys His Ser Ala Ser Pro Gly Ala Gly Arg  *
         35                  40                  45      47


     <210> 1152
     <211> 64
     <212> PRT
     <213> Homo sapiens


     <400> 1152
Met Lys Asp His Leu Glu Phe Pro Phe Leu Asp Leu Leu Asp Leu Thr
  1               5                  10                  15
Asp Ser Leu Gly Leu Leu Gly Phe Gln Gly Leu Leu Ala Leu Leu Ala
             20                  25                  30
Leu Thr Phe Leu Leu Val Met Arg Tyr Val Asn Gln Ala Leu Gln Ala
         35                  40                  45
Pro Gln Asp Leu Gln Val Ile Lys Asp Ser Lys Glu Asn Lys Glu  *
     50                  55                  60          63


     <210> 1153
     <211> 61
     <212> PRT
     <213> Homo sapiens


     <400> 1153
Met Thr Ala Arg Phe Leu Leu Ala Arg Pro Ala Tyr Ser Ser Ala Leu
  1               5                  10                  15
Leu Arg Gly Leu Gly Gly Pro Arg Thr Pro Leu Ile Gln Phe Ser Arg
             20                  25                  30
Cys Gly Met Met Ser Ile Arg Leu Leu Gly Leu Phe Pro Leu Cys Leu
         35                  40                  45
Cys Ser Val Leu Trp Phe Pro Gln Gln His Ser Leu  *
     50                  55                  60


     <210> 1154
     <211> 75
```

<212> PRT
<213> Homo sapiens

<400> 1154
```
Met Asp Ser Thr Phe Leu Ala Thr Arg Ala Val Arg Gly Gln Leu Tyr
 1               5                  10                  15
Leu Trp Ile Ser Met Leu Thr Ile Ala Thr Gly Lys Leu Cys Ala Arg
            20                  25                  30
Cys Tyr Pro Glu Asn Gln Asp His Ile Ile Gln Met Leu Pro Cys Ser
        35                  40                  45
Pro Ala Ser Val Ile Leu His Leu Pro Trp Met Met Lys Phe Phe Leu
    50                  55                  60
Ala Arg His Leu Ile Lys Trp Leu Glu Asn  *
65                  70              74
```

<210> 1155
<211> 68
<212> PRT
<213> Homo sapiens

<400> 1155
```
Met Met Ala Lys Ser Val Arg Phe Cys Tyr Val Leu Phe Val Glu Glu
 1               5                  10                  15
Ile Arg Phe Ala Val Leu Val Val Gln Arg Leu Ala Lys Ser Asp Leu
            20                  25                  30
Trp Ala Lys Ser Gly Leu Leu Ser Ile Phe Ile Phe Ile Ser Lys Val
        35                  40                  45
Leu Leu Lys Gln Thr His Leu Leu Val Cys Arg Met Tyr Ile Ala Ala
    50                  55                  60
Phe Ala Leu  *
65      67
```

<210> 1156
<211> 60
<212> PRT
<213> Homo sapiens

<400> 1156
```
Met Ile Tyr Phe Leu Ser Thr Pro Leu Leu Leu Thr Leu Phe Asn Ile
 1               5                  10                  15
Leu Met Thr Phe Phe Phe Val Ala Pro Pro Leu Asn Leu Leu Asn Lys
            20                  25                  30
Thr His Phe Cys Phe Phe Ser Ser Tyr Ser Leu Lys Asp Phe Arg Cys
        35                  40                  45
Pro Pro Pro Lys Leu Lys Phe Leu Leu His Pro  *
    50                  55              59
```

<210> 1157
<211> 776
<212> PRT

<213> Homo sapiens

<400> 1157
Met Leu Phe Ile Val Thr Ala Leu Leu Cys Cys Gly Leu Cys Asn Gly
1 5 10 15
Val Leu Ile Glu Glu Thr Glu Ile Val Met Pro Thr Pro Lys Pro Glu
20 25 30
Leu Trp Ala Glu Thr Asn Phe Pro Leu Ala Pro Trp Lys Asn Leu Thr
35 40 45
Leu Trp Cys Arg Ser Pro Ser Gly Ser Thr Lys Glu Phe Val Leu Leu
50 55 60
Lys Asp Gly Thr Gly Trp Ile Ala Thr Arg Pro Ala Ser Glu Gln Val
65 70 75 80
Arg Ala Ala Phe Pro Leu Gly Ala Leu Thr Gln Ser His Thr Gly Ser
85 90 95
Tyr His Cys His Ser Trp Glu Glu Met Ala Val Ser Glu Pro Ser Glu
100 105 110
Ala Leu Glu Leu Val Gly Thr Asp Ile Leu Pro Lys Pro Val Ile Ser
115 120 125
Ala Ser Pro Thr Ile Arg Gly Gln Glu Leu Gln Leu Arg Cys Lys Gly
130 135 140
Trp Leu Ala Gly Met Gly Phe Ala Leu Tyr Lys Glu Gly Glu Gln Glu
145 150 155 160
Pro Val Gln Gln Leu Gly Ala Val Gly Arg Glu Ala Phe Phe Thr Ile
165 170 175
Gln Arg Met Glu Asp Lys Asp Glu Gly Asn Tyr Ser Cys Arg Thr His
180 185 190
Thr Glu Lys Arg Pro Phe Lys Trp Ser Glu Pro Ser Glu Pro Leu Glu
195 200 205
Leu Val Ile Lys Glu Met Tyr Pro Lys Pro Phe Phe Lys Thr Trp Ala
210 215 220
Ser Pro Val Val Thr Pro Gly Ala Arg Val Thr Phe Asn Cys Ser Thr
225 230 235 240
Pro His Gln His Met Ser Phe Ile Leu Tyr Lys Asp Gly Ser Glu Ile
245 250 255
Ala Ser Ser Asp Arg Ser Trp Ala Ser Pro Gly Ala Ser Ala Ala His
260 265 270
Phe Leu Ile Ile Ser Val Gly Ile Gly Asp Gly Gly Asn Tyr Ser Cys
275 280 285
Arg Tyr Tyr Asp Phe Ser Ile Trp Ser Glu Pro Ser Asp Pro Val Glu
290 295 300
Leu Val Val Thr Glu Phe Tyr Pro Lys Pro Thr Leu Leu Ala Gln Pro
305 310 315 320
Gly Pro Val Val Phe Pro Gly Lys Ser Val Ile Leu Arg Cys Gln Gly
325 330 335
Thr Phe Gln Gly Met Arg Phe Ala Leu Leu Gln Glu Gly Ala His Val
340 345 350
Pro Leu Gln Phe Arg Ser Val Ser Gly Asn Ser Ala Asp Phe Leu Leu
355 360 365
His Thr Val Gly Ala Glu Asp Ser Gly Asn Tyr Ser Cys Ile Tyr Tyr
370 375 380
Glu Thr Thr Met Ser Asn Arg Gly Ser Tyr Leu Ser Met Pro Leu Met
385 390 395 400
Ile Trp Val Thr Asp Thr Phe Pro Lys Pro Trp Leu Phe Ala Glu Pro
405 410 415
Ser Ser Val Val Pro Met Gly Gln Asn Val Thr Leu Trp Cys Arg Gly
420 425 430
Pro Val His Gly Val Gly Tyr Ile Leu His Lys Glu Gly Glu Ala Thr

```
        435                 440                 445
Ser Met Gln Leu Trp Gly Ser Thr Ser Asn Asp Gly Ala Phe Pro Ile
    450                 455                 460
Thr Asn Ile Ser Gly Thr Ser Met Gly Arg Tyr Ser Cys Cys Tyr His
465                 470                 475                 480
Pro Asp Trp Thr Ser Ser Ile Lys Ile Gln Pro Ser Asn Thr Leu Glu
                485                 490                 495
Leu Leu Val Thr Gly Leu Leu Pro Lys Pro Ser Leu Leu Ala Gln Pro
            500                 505                 510
Gly Pro Met Val Ala Pro Gly Glu Asn Met Thr Leu Gln Cys Gln Gly
        515                 520                 525
Glu Leu Pro Asp Ser Thr Phe Val Leu Leu Lys Glu Gly Ala Gln Glu
    530                 535                 540
Pro Leu Glu Gln Gln Arg Pro Ser Gly Tyr Arg Ala Asp Phe Trp Met
545                 550                 555                 560
Pro Ala Val Arg Gly Glu Asp Ser Gly Ile Tyr Ser Cys Val Tyr Tyr
                565                 570                 575
Leu Asp Ser Thr Pro Phe Ala Ala Ser Asn His Ser Asp Ser Leu Glu
            580                 585                 590
Ile Trp Val Thr Asp Lys Pro Pro Lys Pro Ser Leu Ser Ala Trp Pro
        595                 600                 605
Ser Thr Met Phe Lys Leu Gly Lys Asp Ile Thr Leu Gln Cys Arg Gly
    610                 615                 620
Pro Leu Pro Gly Val Glu Phe Val Leu Glu His Asp Gly Glu Glu Ala
625                 630                 635                 640
Pro Gln Gln Phe Ser Glu Asp Gly Asp Phe Val Ile Asn Asn Val Glu
                645                 650                 655
Gly Lys Gly Ile Gly Asn Tyr Ser Cys Ser Tyr Arg Leu Gln Ala Tyr
            660                 665                 670
Pro Asp Ile Trp Ser Glu Pro Ser Asp Pro Leu Glu Leu Val Gly Ala
        675                 680                 685
Ala Gly Pro Val Ala Gln Glu Cys Thr Val Gly Asn Ile Val Arg Ser
    690                 695                 700
Ser Leu Ile Val Val Val Val Val Ala Leu Gly Val Val Leu Ala Ile
705                 710                 715                 720
Glu Trp Lys Lys Trp Pro Arg Leu Arg Thr Arg Gly Ser Glu Thr Asp
                725                 730                 735
Gly Arg Asp Gln Thr Ile Ala Leu Glu Glu Cys Asn Gln Glu Gly Glu
            740                 745                 750
Pro Gly Thr Pro Ala Asn Ser Pro Ser Ser Thr Ser Gln Arg Ile Ser
        755                 760                 765
Val Glu Leu Pro Val Pro Ile  *
    770                 775
```

<210> 1158
<211> 80
<212> PRT
<213> Homo sapiens

<400> 1158

```
Met Ile Gln Leu Phe Phe Val Leu Tyr Gly Ile Leu Ala Leu Ala Phe
 1               5                  10                  15
Leu Ser Gly Tyr Tyr Val Thr Leu Ala Ala Gln Ile Leu Ala Val Leu
            20                  25                  30
Leu Pro Pro Val Met Leu Leu Ile Asp Gly Asn Val Ala Tyr Trp His
        35                  40                  45
```

Asn Thr Arg Arg Val Glu Phe Trp Asn Gln Met Lys Leu Leu Gly Glu
50 55 60
Ser Val Gly Ile Phe Gly Thr Ala Val Ile Leu Ala Thr Asp Gly *
65 70 75 79

<210> 1159
<211> 132
<212> PRT
<213> Homo sapiens

<400> 1159
Met Ser Ser Gly Thr Glu Leu Leu Trp Pro Gly Ala Ala Leu Leu Val
1 5 10 15
Leu Leu Gly Val Ala Ala Ser Leu Cys Val Arg Cys Ser Arg Pro Gly
20 25 30
Ala Lys Arg Ser Glu Lys Ile Tyr Gln Gln Arg Ser Leu Arg Glu Asp
35 40 45
Gln Gln Ser Phe Thr Gly Ser Arg Thr Tyr Ser Leu Val Gly Gln Ala
50 55 60
Trp Pro Gly Pro Leu Ala Asp Met Ala Pro Thr Arg Lys Asp Lys Leu
65 70 75 80
Leu Gln Phe Tyr Pro Ser Leu Glu Asp Pro Ala Ser Ser Arg Tyr Gln
85 90 95
Asn Phe Ser Lys Gly Ser Arg His Gly Ser Glu Glu Ala Tyr Ile Asp
100 105 110
Pro Thr Ala Ile Lys Tyr Phe Leu Thr Gln Ala Thr Ala Ser Ile Ile
115 120 125
Leu Leu Ile Ala
130 132

<210> 1160
<211> 167
<212> PRT
<213> Homo sapiens

<400> 1160
Met Val Gly Leu Gly Gly Met Ser Gln Leu Leu Leu Ala Ser Leu Leu
1 5 10 15
Pro Pro Val Pro Gln Gly Ser Pro Thr Arg Arg Lys Leu Pro Ala Ser
20 25 30
Leu Leu Val Ser Thr Ala Leu Ile Ser Pro Val Cys Val Arg Gly Trp
35 40 45
Met Trp Gln Asn Leu Gln Asn Arg Ile His Gly Ser His Thr Ser Ala
50 55 60
Arg Arg Val Pro Ser Leu Pro Gly Ala Gly Gln Val Gly Val Arg Trp
65 70 75 80
Glu Ala Gly Pro Ala Cys Arg Thr Gln Pro Ser Pro Gln Asn Leu Ala
85 90 95
Pro Arg Pro His Pro Ser Ala Ala Gln Leu Ile Glu Asn Ala Ala Leu
100 105 110
Arg Ser Ala Met Ser Gly Glu Arg Leu Phe Pro Glu Gly Gln Glu His
115 120 125
Leu Gly Pro Leu Val Ala Pro Arg Val Pro Met Gly Gly Ala Leu Cys

```
    130                 135                 140
Pro Pro Leu Pro Ser Leu Ser Cys Ala Ile Cys Lys Val Gly Ala Ala
145                 150                 155                 160
Arg Glu Ala Gly Gly Arg  *
                165 166
```

<210> 1161
<211> 84
<212> PRT
<213> Homo sapiens

<400> 1161

```
Met Ala Asn Leu Leu Leu Leu Ile Val Pro Ile Leu Ile Ala Met Ala
 1               5                  10                  15
Phe Leu Met Leu Thr Glu Arg Lys Ile Leu Gly Tyr Ile Gln Leu Arg
            20                  25                  30
Lys Gly Pro Asn Val Val Gly Pro Tyr Gly Leu Leu Gln Pro Phe Ala
        35                  40                  45
Asp Ala Ile Lys Leu Phe Thr Lys Glu Pro Leu Lys Pro Ala Thr Ser
    50                  55                  60
Ala Ile Thr Leu Tyr Ile Thr Ala Pro Thr Leu Ala Leu Thr Ile Ala
65                  70                  75                  80
Leu Leu Leu  *
        83
```

<210> 1162
<211> 80
<212> PRT
<213> Homo sapiens

<400> 1162

```
Met Lys Ala Trp Cys Phe Ser Asn Lys Phe Trp Leu Ala Val Leu Pro
 1               5                  10                  15
Ile Cys Cys Ala Ser Ala Ala Tyr Leu Gly Gln Val Trp Leu Leu Ile
            20                  25                  30
Tyr Ala Trp Arg Ala Glu Thr Ser Leu Glu Thr Glu Phe Tyr Thr Ile
        35                  40                  45
Pro Leu Ser Trp Leu Tyr Tyr Phe Thr Thr Thr Tyr Tyr Leu Met Phe
    50                  55                  60
Leu Pro Ser Leu Lys Phe Ala Gln Asp Ser Pro Pro Arg Ala Phe  *
65                  70                  75                  79
```

<210> 1163
<211> 71
<212> PRT
<213> Homo sapiens

<400> 1163

```
Met Tyr Gly Leu Lys Ile Leu Ser His Leu Trp Val Leu Leu Ile Leu
 1               5                  10                  15
```

Ser Leu Leu Leu Phe Leu Arg Lys Ser Phe Lys Phe Tyr Ala Val Ser
20 25 30
Phe Val Cys Phe Ala Phe Val Ala Phe Trp Asn Asn Leu Gln Lys Ile
35 40 45
Ile Ala Gln Ala Asn Val Ile Gln Ser Pro Ser Ile Phe Pro Cys Ser
50 55 60
Ser Ser Thr Phe Lys Leu *
65 70

<210> 1164
<211> 56
<212> PRT
<213> Homo sapiens

<400> 1164
Met Glu Thr Ala Val Ile Gly Val Val Val Val Leu Phe Val Val Thr
1 5 10 15
Val Ala Ile Thr Cys Val Leu Cys Cys Phe Ser Cys Asp Ser Arg Ala
20 25 30
Gln Asp Pro Gln Gly Gly Pro Gly Arg Ser Phe Thr Val Ala Thr Phe
35 40 45
Arg Gln Glu Ala Ser Leu Phe Thr
50 55 56

<210> 1165
<211> 97
<212> PRT
<213> Homo sapiens

<221> misc_feature
<222> (1)...(97)
<223> Xaa = any amino acid or nothing

<400> 1165
Met Lys Met Leu Cys Gly Leu Leu Arg Thr Val Gln Gly Val Arg Phe
1 5 10 15
Pro Gln Leu Thr Arg Ile His Gly Pro Ser Thr Gln Gly His Gln Leu
20 25 30
Leu Leu Leu Trp Val Gly Val Leu Gln Val Gly Xaa Ser Ser Leu Gly
35 40 45
Leu Gln Asn Asp Leu Met Gly Pro Ser Leu Gly Arg Gly Pro Pro Pro
50 55 60
Leu Ala Ala Ser Thr Arg Cys Arg His Val Ala Gln Leu Gly Val Gly
65 70 75 80
Leu Ser Lys Thr Trp Gln Pro Ser Thr His Gly Ile Ala Ser Ala Pro
85 90 95 96
*

<210> 1166
<211> 48

<212> PRT
<213> Homo sapiens

<400> 1166
Met Leu Ile Phe Val Phe Leu Phe Ser Tyr Leu Ile Ala Leu Ala Gly
1 5 10 15
Thr Phe Ser Pro Arg Leu Asn Arg Ser Gly Glu Ser Val His Pro Phe
20 25 30
Ala Leu His Pro Val Leu Arg Arg Lys His Pro Val Ile His Leu *
35 40 45 47

<210> 1167
<211> 274
<212> PRT
<213> Homo sapiens

<400> 1167
Met Glu Ala Pro Leu Ser His Leu Glu Ser Arg Tyr Leu Pro Ala His
1 5 10 15
Phe Ser Pro Leu Val Phe Phe Leu Leu Leu Ser Ile Met Met Ala Cys
20 25 30
Cys Leu Val Ala Phe Phe Val Leu Gln Arg Gln Pro Arg Cys Trp Glu
35 40 45
Ala Ser Val Glu Asp Leu Leu Asn Asp Gln Val Thr Leu His Ser Ile
50 55 60
Arg Pro Arg Glu Glu Asn Asp Leu Gly Pro Ala Gly Thr Val Asp Ser
65 70 75 80
Ser Gln Gly Gln Gly Tyr Leu Glu Glu Lys Ala Ala Pro Cys Cys Pro
85 90 95
Ala His Leu Ala Phe Ile Tyr Thr Leu Val Ala Phe Val Asn Ala Leu
100 105 110
Thr Asn Gly Met Leu Pro Ser Val Gln Thr Tyr Ser Cys Leu Ser Tyr
115 120 125
Gly Pro Val Ala Tyr His Leu Ala Ala Thr Leu Ser Ile Val Ala Asn
130 135 140
Pro Leu Ala Ser Leu Val Ser Met Phe Leu Pro Asn Arg Ser Leu Leu
145 150 155 160
Phe Leu Gly Val Leu Ser Val Leu Gly Thr Cys Phe Gly Gly Tyr Asn
165 170 175
Met Ala Met Ala Val Met Ser Pro Cys Pro Leu Leu Gln Gly His Trp
180 185 190
Gly Gly Glu Val Leu Ile Val Ser Ile Arg Pro Val Ala Ser Trp Val
195 200 205
Leu Phe Ser Gly Cys Leu Ser Tyr Val Lys Val Met Leu Gly Val Val
210 215 220
Leu Arg Asp Leu Ser Arg Ser Ala Leu Leu Trp Cys Gly Ala Ala Val
225 230 235 240
Gln Leu Gly Ser Leu Leu Gly Ala Leu Leu Met Phe Pro Leu Val Asn
245 250 255
Val Leu Arg Leu Phe Ser Ser Ala Asp Phe Cys Asn Leu His Cys Pro
260 265 270
Ala *
273

<210> 1168
<211> 230
<212> PRT
<213> Homo sapiens

<400> 1168
Met Arg Ile Cys Asn Leu Ile Ser Met Met Leu Leu Leu Cys His Trp
1               5                   10                  15
Asp Gly Cys Leu Gln Phe Leu Val Pro Met Leu Gln Asp Phe Pro Arg
            20                  25                  30
Asn Cys Trp Val Ser Ile Asn Gly Met Val Asn His Ser Trp Ser Glu
        35                  40                  45
Leu Tyr Ser Phe Ala Leu Phe Lys Ala Met Ser His Met Leu Cys Ile
    50                  55                  60
Gly Tyr Gly Arg Gln Ala Pro Glu Ser Met Thr Asp Ile Trp Leu Thr
65                  70                  75                  80
Met Leu Ser Met Ile Val Gly Ala Thr Cys Tyr Ala Met Phe Ile Gly
                85                  90                  95
His Ala Thr Ala Leu Ile Gln Ser Leu Asp Ser Ser Arg Arg Gln Tyr
            100                 105                 110
Gln Glu Lys Tyr Lys Gln Val Glu Gln Tyr Met Ser Phe His Lys Leu
        115                 120                 125
Pro Ala Asp Phe Arg Gln Lys Ile His Asp Tyr Tyr Glu His Arg Tyr
    130                 135                 140
Gln Gly Lys Met Phe Asp Glu Asp Ser Ile Leu Gly Glu Leu Asn Gly
145                 150                 155                 160
Pro Leu Arg Glu Glu Ile Val Asn Phe Asn Cys Arg Lys Leu Val Ala
                165                 170                 175
Ser Met Pro Leu Phe Ala Asn Ala Asp Pro Asn Phe Val Thr Ala Met
            180                 185                 190
Leu Thr Lys Leu Lys Phe Glu Val Phe Gln Pro Gly Asp Tyr Ile Ile
        195                 200                 205
Pro Arg Arg His His Arg Glu Glu Asp Val Leu His Pro Ala Arg Arg
    210                 215                 220
Gly Gln Arg Ala His *
225             229

<210> 1169
<211> 213
<212> PRT
<213> Homo sapiens

<400> 1169
Met Ala His Phe Thr Trp Ala His Leu Arg Val Leu Thr Leu Phe Leu
1               5                   10                  15
Leu Gln Val Gly Leu Leu Asp Asp Val His Gln Leu Leu Gly Pro Gln
            20                  25                  30
Ala Asp Glu Asp Ser Leu Ser Ile Phe Thr Val Met Pro Ala Leu His
        35                  40                  45
Gln Ser Gln Glu Gln Leu Gly Gly Ile Val Leu Glu Leu Gln His Gln
    50                  55                  60
Ile His Ala Val Leu Ala Gln Gly Ala Asp Val Ile Glu Asp Gln Cys
65                  70                  75                  80
Gly Asp Asp Val Tyr Ala Ile Gly Leu Val Ser His Asn Ala Ser Leu

85 90 95
Val Leu Met Ala Gly Ala Leu Ala Val Leu Ser Glu Gly Leu Gln Gly
100 105 110
Leu Asp Asp Glu Ala His Val Val Leu Ile Asp Val Glu Pro Gln Gln
115 120 125
Pro Gln Ala Ala Arg Gly Ala Ala Ala His Asp Val Gln Glu Leu Gln
130 135 140
Arg Leu Ala Tyr Gln Val Val Val Gly Phe Val Val Leu Thr Ala Gln
145 150 155 160
Glu Val Leu Gln Val Pro Val Val Val Leu Thr Gln Gln Leu Gln Lys
165 170 175
Ala Gln Asp Gly Leu His Asp Glu His Gly Cys Ala His Leu Thr Ala
180 185 190
Leu His Thr Phe Ala His Leu Val Pro Pro Ala Gln Ala Gly Ala Gln
195 200 205
Arg Val Ala Gly *
210 212

<210> 1170
<211> 51
<212> PRT
<213> Homo sapiens

<400> 1170
Met Tyr Ser Leu Val Leu Thr Phe Leu Val Ser Phe Cys Ala Leu Ser
1 5 10 15
Lys Thr Phe Leu Asp His Trp Phe Gln Met Phe Ile Tyr Tyr Ile Leu
20 25 30
Phe Lys Asp Ser Glu Ile Gly Phe Cys His Pro Leu Leu Tyr Val Leu
35 40 45
Phe His *
50

<210> 1171
<211> 157
<212> PRT
<213> Homo sapiens

<400> 1171
Met Leu Val Pro Leu Asn Leu Cys Leu Gln Ser Thr Leu Ala Leu Val
1 5 10 15
Ser Leu Pro Leu Pro Gly Ile Gly Arg Ala Phe Cys Glu Trp Leu Ser
20 25 30
Gly Thr Phe Lys Ala Arg Arg Gln Gly Pro Lys Ala Lys Arg Glu Leu
35 40 45
Trp Asp Val Pro Ser Pro Val Arg Gly Trp Pro Trp Gly Phe Arg Leu
50 55 60
Arg Gly Val Pro Gly Pro Val Ser Pro Ala Phe Gly Pro Phe Gly Glu
65 70 75 80
Phe Gly Glu Glu Val Pro Thr Ala Arg Pro Gly Asp Val Arg Gly Ala
85 90 95
Ala Leu Thr Phe Ile Val Gly Val Ser Ser Glu Val Ser Val Gln Arg
100 105 110

Arg Ser Ala Gly Arg Ser His Arg Gly Arg Arg Arg Arg Ala Ser Cys
115 120 125
Thr Ala Ala Pro Gly Gly Gly Val Thr Arg Arg Trp Lys Glu Tyr Cys
130 135 140
Thr Gln Arg Ile Asn Asn Leu Val Lys Pro Phe Ser *
145 150 155 156

<210> 1172
<211> 69
<212> PRT
<213> Homo sapiens

<400> 1172
Met Asn Pro Tyr Ile Ser Ile Ile Val Phe Ile Val Phe Leu Cys Ser
1 5 10 15
Glu Asn Tyr Pro Trp Asn Asn Met Leu Arg Ile Thr Gly Ser Ser Pro
20 25 30
Tyr Leu His Phe Leu Ser Val Leu Gly Val Leu Val Asn Ser Tyr Val
35 40 45
Leu Ile Leu Phe Asn Ser Glu Phe Leu Thr Gln His Phe Arg Glu Arg
50 55 60
Ile Gln Ala Gly *
65 68

<210> 1173
<211> 75
<212> PRT
<213> Homo sapiens

<400> 1173
Met Cys Ser Leu Lys Phe Trp Ile Cys Phe Cys Gln Ala Val Ser Met
1 5 10 15
His Leu Cys Ala Thr Gln Leu Ser Val Ser Leu Pro Ala Gly Ile Ser
20 25 30
Met Phe Val Ser Gly Leu Val Cys Asp Ile Cys Val Trp Ser Gly Ser
35 40 45
Gly Met Thr His Pro Tyr Trp Ser Arg Met Arg Val Glu Met Met Val
50 55 60
Ala Gly Cys Phe Arg Glu Arg Asp Ala His *
65 70 74

<210> 1174
<211> 77
<212> PRT
<213> Homo sapiens

<400> 1174
Met Leu Ser Ser Phe Phe Lys Ser Cys Phe Cys Val Ser Phe Trp Thr
1 5 10 15
Leu Ser Ile Ala Thr Ser Ser Asn Leu Leu Ile Phe Ser Ser Ala Ile

```
            20                      25                      30
Ser Asn Leu Leu Leu Ile Leu Ser Ser Val Phe Ser Ile Leu Asp Ile
        35                      40                      45
Val Val Phe Ile Thr Arg Ser Met Ile Trp Phe Cys Phe His Pro Cys
    50                      55                      60
Ile Tyr Ile Thr Cys Pro Val Phe His Ser Ala Ser  *
 65                      70                      75  76
```

<210> 1175
<211> 59
<212> PRT
<213> Homo sapiens

<400> 1175

```
Met Ser Phe Ala Phe Ser Leu Trp Tyr Pro Phe Leu Arg Asp Leu Arg
  1               5                  10                  15
Ser Cys Phe Lys Leu Ser Lys Leu Ser Cys His Ser Pro Ile Ser Phe
            20                  25                  30
Val Gln Tyr Thr Thr Met Ser Thr Arg Val Ser Cys Leu Asn Leu Leu
        35                  40                  45
Tyr Pro His Leu Arg Val Val Ser Ile His Ser
    50                  55              59
```

<210> 1176
<211> 55
<212> PRT
<213> Homo sapiens

<400> 1176

```
Met His Leu Leu Cys Ser Gly His Lys Leu Cys Leu Cys Ile Val Tyr
  1               5                  10                  15
Ile Ser Phe Phe Leu Phe Phe Lys Val Tyr Gly Phe Cys Phe Leu His
            20                  25                  30
Ala Asn Ile Val Asn Tyr Thr Glu Asp Thr Thr Asp Ser Ile Tyr Lys
        35                  40                  45
Val Tyr Arg Asn Ile Ile  *
    50              54
```

<210> 1177
<211> 86
<212> PRT
<213> Homo sapiens

<400> 1177

```
Met Leu Ser Met Leu Leu Arg Ala Val Phe Cys Cys Cys Arg Arg Leu
  1               5                  10                  15
His Leu Val Ser Ser Ile Leu Phe Cys Cys Ser Arg Asn Arg Thr Leu
            20                  25                  30
Ser Met Lys Glu Ala Asn Leu Leu Leu Arg Val Leu Ile Cys Ser Phe
        35                  40                  45
```

```
Ser Trp Val Arg Thr Ala Trp Met Leu Gly Ser Thr Ser Arg Thr Arg
     50                  55                  60
Gly Leu Ser Arg Leu Trp Leu Thr Val Thr Ala Val Met Pro Pro Met
 65                  70                  75                  80
Pro Leu Ala Pro Pro  *
                 85


     <210> 1178
     <211> 189
     <212> PRT
     <213> Homo sapiens


     <400> 1178
Met Met Pro Leu Leu Ser Leu Ile Phe Ser Ala Leu Phe Ile Leu Phe
  1               5                  10                  15
Gly Thr Val Ile Val Gln Ala Phe Ser Asp Ser Asn Asp Glu Arg Glu
             20                  25                  30
Ser Ser Pro Pro Glu Lys Glu Glu Ala Gln Glu Lys Thr Gly Lys Thr
         35                  40                  45
Glu Pro Ser Phe Thr Lys Glu Asn Ser Ser Lys Ile Pro Lys Lys Gly
     50                  55                  60
Phe Val Glu Val Thr Glu Leu Thr Asp Val Thr Tyr Thr Ser Asn Leu
 65                  70                  75                  80
Val Arg Leu Arg Pro Gly His Met Asn Val Val Leu Ile Leu Ser Asn
                 85                  90                  95
Ser Thr Lys Thr Ser Leu Leu Gln Lys Phe Ala Leu Glu Val Tyr Thr
            100                 105                 110
Phe Thr Gly Ser Ser Cys Leu His Phe Ser Phe Leu Ser Leu Asp Lys
        115                 120                 125
His Arg Glu Trp Leu Glu Tyr Leu Leu Glu Phe Ala Gln Asp Ala Ala
    130                 135                 140
Pro Ile Pro Asn Gln Tyr Asp Lys His Phe Met Glu Arg Asp Tyr Thr
145                 150                 155                 160
Gly Tyr Val Leu Ala Leu Asn Gly His Lys Lys Tyr Phe Cys Leu Phe
                165                 170                 175
Lys Pro Gln Lys Thr Val Glu Glu Gly Gly Lys Pro  *
            180                 185         188


     <210> 1179
     <211> 55
     <212> PRT
     <213> Homo sapiens


     <400> 1179
Met Ile Cys Lys Tyr Phe Phe Leu Ile Leu Trp Val Val Phe Ser Phe
  1               5                  10                  15
Phe Phe Met Phe Leu Asp Ala Gln Lys Phe Ile Ile Leu Met Lys Ser
             20                  25                  30
Asn Ser Ser Phe Leu Leu Leu Leu His Met Leu Leu Glu Ser Tyr Leu
         35                  40                  45
Arg Asn His Cys Gln Ile  *
     50              54
```

<210> 1180
<211> 81
<212> PRT
<213> Homo sapiens

<400> 1180
Met Ala Phe Leu Leu Ser Thr Leu Leu Asn His Tyr Leu Ala Cys Lys
1 5 10 15
His Ser Ser Glu Leu Trp Leu Gln Ser Ser Leu Asn Asn Leu Gly Lys
20 25 30
Lys Lys Asp Lys Ala Tyr Ile Phe Thr Val Leu Ala Leu Lys His Ile
35 40 45
Pro Gln Met Pro Leu Arg Ile Tyr Phe Val Leu Gly Gln Ser Trp Trp
50 55 60
Leu Met Pro Val Ile Pro Ala Ile Trp Glu Ala Glu Ala Arg Thr Ala
65 70 75 80
*

<210> 1181
<211> 69
<212> PRT
<213> Homo sapiens

<400> 1181
Met Asp Glu Val His Val Leu Gly Leu Ala Leu Leu Thr Val Leu Ile
1 5 10 15
Glu Leu Val Ser Pro Leu Asp Ser Leu Arg Arg His Ser Cys Tyr Ile
20 25 30
Thr His Thr Phe Ser Cys Asn His Thr Asn Ser His Phe Tyr Ile Leu
35 40 45
Ser Ile Ser Cys Thr Asn Trp Gly Leu Lys Val Tyr Lys Ile Phe Leu
50 55 60
Ser Cys Glu Phe *
65 68

<210> 1182
<211> 430
<212> PRT
<213> Homo sapiens

<400> 1182
Met Ile Thr Lys Thr Pro Ala Gln Leu Arg Ser Val Ala Thr Ile Leu
1 5 10 15
Lys Thr Leu Cys Leu Ala Ser Pro Thr Val Ala Asn Val Lys Ala Pro
20 25 30
Pro Gln Val Ala Val Ala Ala Gly Thr Pro Asn Thr Ser Gly Ser Ile
35 40 45
His Glu Asn Pro Pro Lys Ala Lys Ala Thr Val Asn Val Lys Gln Ala
50 55 60

```
Ala Lys Val Val Lys Ala Ser Ser Pro Ser Tyr Leu Ala Glu Gly Lys
 65                  70                  75                  80
Ile Arg Cys Leu Ala Gln Pro His Pro Gly Thr Gly Val Pro Arg Ala
                 85                  90                  95
Ala Ala Glu Leu Pro Leu Glu Ala Glu Lys Ile Lys Thr Gly Thr Gln
            100                 105                 110
Lys Gln Ala Lys Thr Asp Met Ala Phe Lys Thr Ser Val Ala Val Glu
        115                 120                 125
Met Ala Gly Ala Pro Ser Trp Thr Lys Val Ala Glu Glu Gly Asp Lys
    130                 135                 140
Pro Pro His Gly Pro Arg Cys Pro Asn His Ala Cys Gln Arg Leu Gly
145                 150                 155                 160
Gly Leu Ser Ala Pro Pro Trp Ala Lys Pro Glu Asp Arg Gln Thr Gln
                165                 170                 175
Pro Gln Pro His Gly His Val Pro Gly Lys Thr Thr Gln Gly Gly Pro
            180                 185                 190
Cys Pro Ala Ala Cys Glu Val Gln Gly Met Leu Val Pro Pro Met Ala
        195                 200                 205
Pro Thr Gly His Ser Thr Cys Asn Val Glu Ser Trp Gly Asp Asn Gly
    210                 215                 220
Ala Thr Arg Ala Gln Pro Ser Met Pro Gly Gln Ala Val Pro Cys Gln
225                 230                 235                 240
Glu Asp Thr Val Gly Ser Leu Leu Ala Ser Leu Cys Ala Glu Val Ala
                245                 250                 255
Gly Val Leu Ala Ser Gln Glu Asp Leu Arg Thr Leu Leu Ala Lys Ala
            260                 265                 270
Leu Ser Gln Gly Glu Val Trp Ala Ala Leu Asn Gln Ala Leu Ser Lys
        275                 280                 285
Glu Val Leu Gly Ala Thr Val Thr Lys Ala Leu Pro Gln Ser Met Leu
    290                 295                 300
Ser Met Ala Leu Val Lys Ala Leu Ser Trp Ser Glu Leu Arg Leu Thr
305                 310                 315                 320
Leu Ser Arg Ala Leu Ser Arg Gly Glu Leu Arg Ala Glu Leu Thr Lys
                325                 330                 335
Val Met Gln Gly Lys Leu Ala Glu Val Leu Ser Lys Ala Leu Thr Glu
            340                 345                 350
Glu Glu Trp Val Ala Leu Ser Gln Ala Leu Cys Gln Gly Glu Leu Gly
        355                 360                 365
Ala Leu Leu Ser Gln Ser Trp Cys Arg Val Ala Leu Arg Thr Gly Thr
    370                 375                 380
Ile Leu Pro Lys Ala Ala Ser Lys Ser Thr Gly Ser Gly Val Thr Lys
385                 390                 395                 400
Thr Pro Ala Leu Val Lys Val Ala Cys Arg Arg Ser Pro Ser Ala Ala
                405                 410                 415
Trp Gly Pro Ser Leu Gly Pro Val Arg Pro Gln Thr Ser Lys
            420                 425                 430
```

<210> 1183
<211> 53
<212> PRT
<213> Homo sapiens

<400> 1183

```
Met Thr Phe Ile Leu Ser Arg Pro Pro Phe Phe Phe Leu Phe Ser Lys
  1               5                  10                  15
Arg Ser Cys Ser Gly Ala Arg Trp Ser Arg Trp Pro Gln Phe Gly Tyr
```

20 25 30
Ser Thr Ser Pro Pro Gly Ser Met Phe Phe Ser Ser Pro Pro Ser Arg
35 40 45
Gly Ile Pro Ala *
50 52

<210> 1184
<211> 56
<212> PRT
<213> Homo sapiens

<400> 1184
Met Ser Met Leu His Trp Ile His Phe Ile Leu His Val Ser Ile Val
1 5 10 15
Leu Lys Phe Leu Ser Val Lys Cys Ser Ile Ile Tyr Lys Lys Ser Phe
20 25 30
Ala Ser Ser Ala Phe Phe Leu Val Gln Ala Ser Phe Phe His Ile Met
35 40 45
Leu Ser Gln Leu Tyr Phe Gln *
50 55

<210> 1185
<211> 294
<212> PRT
<213> Homo sapiens

<400> 1185
Met Pro Tyr Val Thr Glu Ala Thr Arg Val Gln Leu Val Leu Pro Leu
1 5 10 15
Leu Val Ala Glu Ala Ala Ala Ala Pro Ala Phe Leu Glu Ala Phe Ala
20 25 30
Ala Asn Val Leu Glu Pro Arg Glu His Ala Leu Leu Thr Leu Leu Leu
35 40 45
Val Tyr Gly Pro Arg Glu Gly Gly Arg Gly Ala Pro Asp Pro Phe Leu
50 55 60
Gly Val Lys Ala Ala Ala Ala Glu Leu Glu Arg Arg Tyr Pro Gly Thr
65 70 75 80
Arg Leu Ala Trp Leu Ala Val Arg Ala Glu Ala Pro Ser Gln Val Arg
85 90 95
Leu Met Asp Val Val Ser Lys Lys His Pro Val Asp Thr Leu Phe Phe
100 105 110
Leu Thr Thr Val Trp Thr Arg Pro Gly Pro Glu Val Leu Asn Arg Cys
115 120 125
Arg Met Asn Ala Ile Ser Gly Trp Gln Ala Phe Phe Pro Val His Phe
130 135 140
Gln Glu Phe Asn Pro Ala Leu Ser Pro Gln Arg Ser Pro Pro Gly Pro
145 150 155 160
Pro Gly Ala Gly Pro Asp Pro Pro Ser Pro Pro Gly Ala Asp Pro Ser
165 170 175
Arg Gly Ala Pro Ile Gly Gly Arg Phe Asp Arg Gln Ala Ser Ala Glu
180 185 190
Gly Cys Phe Tyr Asn Ala Asp Tyr Leu Ala Ala Arg Ala Arg Leu Ala
195 200 205

```
Gly Glu Leu Ala Gly Gln Glu Glu Glu Glu Ala Leu Glu Gly Leu Glu
    210                 215                 220
Val Met Asp Val Phe Leu Arg Phe Ser Gly Leu His Leu Phe Arg Ala
225                 230                 235                 240
Val Glu Pro Gly Leu Val Gln Lys Phe Ser Leu Arg Asp Cys Ser Pro
                245                 250                 255
Arg Leu Ser Glu Glu Leu Tyr His Arg Cys Arg Leu Ser Asn Leu Glu
            260                 265                 270
Gly Leu Gly Gly Arg Ala Gln Leu Ala Met Ala Leu Phe Glu Gln Glu
        275                 280                 285
Gln Ala Asn Ser Thr  *
    290         293
```

<210> 1186
<211> 57
<212> PRT
<213> Homo sapiens

<400> 1186

```
Met Met Tyr Ile Leu Leu Val Phe Leu Thr Leu Trp Leu Leu Ile Glu
  1               5                  10                  15
Met Ile His Cys Leu Gln Asn Gly Asp His Arg Arg Thr Arg Pro Pro
             20                  25                  30
Thr Glu Thr Gly Trp Leu Pro Leu Arg Phe His Leu Arg Thr Gly Lys
         35                  40                  45
Ile Leu Arg Tyr Leu Arg Gly Glu  *
     50                  55  56
```

<210> 1187
<211> 191
<212> PRT
<213> Homo sapiens

<400> 1187

```
Met Asp Leu Asp Asn Ala Lys Tyr Ser Leu Leu Gly Phe Ala Leu Phe
  1               5                  10                  15
Trp Val Val Val Gly Phe Phe Phe Val Cys Leu Phe Trp Phe Leu Val
             20                  25                  30
Phe Leu Pro Trp Cys Lys Thr Val Glu Ser Cys Leu Phe Thr Gly Leu
         35                  40                  45
Gly Ser Ile Glu Val Cys Val Ser Ser Val Arg Phe Leu Leu Arg Thr
     50                  55                  60
Ile Cys Ile Phe Asn Asn Ser Thr Ser Ser Arg Pro Ser Arg Arg Asn
 65                  70                  75                  80
Glu Arg Gly Leu Val Ser Ser Pro Glu Leu Ala Leu Glu Cys Val His
                 85                  90                  95
Leu Ala Ala His Gly Leu Val Ala Leu Arg Gly Leu Ile Gln Leu Pro
            100                 105                 110
Leu Gln Leu Pro Ala Val Gly Val Asp Ala Leu Gly Leu Leu Leu Cys
        115                 120                 125
Leu Leu Gln Leu Pro Leu Glu Leu Leu Asp Pro Gly Ile Ala Phe Leu
    130                 135                 140
Cys Leu Leu Leu Val Leu Leu Gly His Leu Ala Leu Val Leu His Leu
```

```
145                 150                 155                 160
Gln Gln Asp Phe Leu Gln Leu Leu Val Phe Leu Leu Gln Arg Leu Gly
                165                 170                 175
Gly Arg Leu Phe Leu Ser Gly Leu Leu Leu Asp Leu Leu Leu  *
            180                 185                 190
```

<210> 1188
<211> 216
<212> PRT
<213> Homo sapiens

<400> 1188

```
Met Ser Pro Pro Leu Leu Leu Leu Pro Leu Leu Leu Leu Leu Pro Leu
 1               5                  10                  15
Leu Asn Val Glu Pro Ala Gly Ala Thr Leu Ile Arg Ile Pro Leu Arg
            20                  25                  30
Gln Val His Pro Gly Arg Arg Thr Leu Asn Leu Leu Arg Gly Trp Gly
        35                  40                  45
Lys Pro Ala Glu Leu Pro Lys Leu Gly Ala Pro Ser Pro Gly Asp Lys
    50                  55                  60
Pro Ala Ser Val Pro Leu Ser Lys Phe Leu Asp Ala Gln Tyr Phe Gly
65                  70                  75                  80
Glu Ile Gly Leu Gly Thr Pro Pro Gln Asn Phe Thr Val Ala Phe Asp
                85                  90                  95
Thr Gly Ser Ser Asn Leu Trp Val Pro Ser Arg Arg Cys His Phe Phe
            100                 105                 110
Ser Val Pro Cys Trp Phe His His Arg Phe Asn Pro Asn Ala Ser Ser
        115                 120                 125
Ser Phe Lys Pro Ser Gly Thr Lys Phe Ala Ile Gln Tyr Gly Thr Gly
    130                 135                 140
Arg Val Asp Gly Ile Leu Ser Glu Asp Lys Leu Thr Ile Gly Gly Ile
145                 150                 155                 160
Lys Gly Ala Ser Val Ile Phe Gly Glu Ala Leu Trp Gly Ile Gln Pro
                165                 170                 175
Gly Ser Ser Leu Phe Pro Ala Pro Met Gly Tyr Trp Gly Leu Gly Phe
            180                 185                 190
Pro Ile Leu Val Leu Trp Glu Gly Ile Ser Ala Pro Ala Gly Cys Thr
        195                 200                 205
Gly Gly Ala Gly Ala Ile Gly  *
    210                 215
```

<210> 1189
<211> 176
<212> PRT
<213> Homo sapiens

<400> 1189

```
Met Ala Leu Arg Gly Ala Leu Gln Ser Gln Ser Gly Leu Leu Ser Leu
 1               5                  10                  15
Leu Leu Leu Gly Leu Gly Asp Lys Asp Pro Val Val Arg Cys Ser Ala
            20                  25                  30
Ser Phe Ala Val Gly Asn Ala Ala Tyr Gln Ala Gly Pro Leu Gly Pro
        35                  40                  45
```

Ala Leu Ala Ala Ala Val Pro Ser Met Thr Gln Leu Leu Gly Asp Pro
50 55 60
Gln Ala Gly Ile Arg Arg Asn Val Ala Ser Ala Leu Gly Asn Leu Gly
65 70 75 80
Pro Glu Gly Leu Gly Glu Glu Leu Leu Gln Cys Glu Val Pro Gln Arg
85 90 95
Leu Leu Glu Met Ala Cys Gly Asp Pro Gln Pro Asn Val Lys Glu Ala
100 105 110
Ala Leu Ile Ala Leu Arg Ser Leu Gln Gln Glu Pro Gly Ile His Gln
115 120 125
Val Leu Val Ser Leu Gly Ala Ser Glu Lys Leu Ser Leu Leu Ser Leu
130 135 140
Gly Asn Gln Ser Leu Pro His Ser Ser Pro Arg Pro Ala Ser Ala Lys
145 150 155 160
His Cys Arg Lys Leu Ile His Leu Leu Arg Pro Ala His Ser Met *
165 170 175

<210> 1190
<211> 58
<212> PRT
<213> Homo sapiens

<400> 1190
Met Ala Gly Thr Ala Gln Leu Leu Gly Leu Lys Gln Leu Ile Gly Leu
1 5 10 15
Glu Leu Leu Thr Ala Gln Cys Gly Gln Ile Thr Gly Tyr Arg Asp Arg
20 25 30
Arg Glu Glu Leu Leu Pro Pro Arg Phe Leu Ala Thr Gly Pro Pro Ser
35 40 45
Cys His Pro Pro Ser Gln Thr Val Pro *
50 55 57

<210> 1191
<211> 88
<212> PRT
<213> Homo sapiens

<400> 1191
Met Gly Ile Cys Leu Thr Trp Lys Pro Pro Thr Gly Val Ser Val Ile
1 5 10 15
Leu Ile Leu Leu Ser Glu Leu His Met Lys Ser Pro Gly Arg Leu Lys
20 25 30
Pro Lys Ser Ser Pro His Phe Ser Thr Val Leu Thr Pro Leu Thr Phe
35 40 45
Met Tyr Pro Gly Leu Ala Leu Leu His Ser Leu Tyr Trp His Trp Gln
50 55 60
Glu Asn Gly Glu Ile Leu Cys Arg Ala Ala Glu Pro Lys Phe Ala Gln
65 70 75 80
Glu Ser Lys Cys Thr Ile Tyr *
85 87

<210> 1192
<211> 136
<212> PRT
<213> Homo sapiens

<400> 1192
Met Val Cys Leu Arg Leu Pro Gly Gly Ser Cys Met Ala Val Leu Thr
1 5 10 15
Val Thr Leu Met Val Leu Ser Ser Pro Leu Ala Leu Ala Gly Asp Thr
20 25 30
Arg Pro Arg Phe Leu Glu Tyr Ser Thr Ser Glu Cys His Phe Phe Asn
35 40 45
Gly Thr Glu Arg Val Arg Tyr Leu Asp Arg Tyr Phe His Asn Gln Glu
50 55 60
Glu Asn Val Arg Phe Asp Ser Asp Val Gly Glu Phe Arg Ala Val Thr
65 70 75 80
Glu Leu Gly Arg Pro Asp Ala Glu Tyr Trp Asn Ser Gln Lys Asp Leu
85 90 95
Leu Gly Thr Ala Arg Arg Thr Ser Trp Ser Arg Ser Gly Ala Gly Trp
100 105 110
Thr Thr Thr Ala Asp Thr Thr Thr Gly Leu Trp Arg Ala Ser Gln Cys
115 120 125
Ser Gly Glu Ser Ile Leu Arg *
130 135

<210> 1193
<211> 99
<212> PRT
<213> Homo sapiens

<400> 1193
Met Leu Ala Ser Arg Gln Ala Cys Cys Pro Pro Val Ser Ser Leu Phe
1 5 10 15
Leu Pro Leu Ser Pro Thr Leu Ser Gly Phe Phe Thr Val Cys Ser Val
20 25 30
Ser His Leu His Val Pro Arg Gly Pro Ala Arg Leu Cys Pro Arg Met
35 40 45
Ser His Gly Ser Pro Ser Gly Leu Pro Ala Glu Pro Ser Glu His Gly
50 55 60
Cys Leu Leu Val Val Gly Leu Gln Gln Asn Cys Thr Arg Leu Thr Ser
65 70 75 80
Pro Ile Leu Ser Ser Arg Gly Leu Arg Val Gln Arg Arg Val Asn Leu
85 90 95
Ala Asp *
98

<210> 1194
<211> 50
<212> PRT
<213> Homo sapiens

<400> 1194

Met Phe Ser Pro Ser Phe Gln Gly Ile Ile Thr Lys Val Arg Cys Val
1 5 10 15
Cys Val Ser Leu Ser Leu Cys Val Cys Val Cys Val Cys Val Cys Val
20 25 30
Cys Val Tyr Lys Glu Pro Gly Met Arg Ala Gly Arg Gly Gly Ser Arg
35 40 45
Leu *
49

<210> 1195
<211> 58
<212> PRT
<213> Homo sapiens

<400> 1195
Met Gln Gly Val Arg Val Ser Phe Gly Trp Ala Met Gly Leu Ala Trp
1 5 10 15
Gly Ser Cys Ala Leu Glu Ala Phe Ser Gly Thr Leu Leu Leu Ser Ala
20 25 30
Ala Trp Thr Leu Ser Leu Ser Pro Pro Ile Cys Gly His Leu Ser Pro
35 40 45
Gln Gln Val Gly Gly Arg Gly Gly Asp *
50 55 57

<210> 1196
<211> 132
<212> PRT
<213> Homo sapiens

<400> 1196
Met Leu Pro Asn Ser Ser Ser Leu Trp Leu Val Met Arg Ile Leu Ile
1 5 10 15
Phe Cys Val Ile Pro Ala Gly Gly Val Leu Gly Ala Pro Thr Ala Ala
20 25 30
Gly Leu Arg Pro Thr Gly Asp Val Ala Leu Arg Arg Pro Ala Gly Ser
35 40 45
Val Glu Pro Ser Gly Ser Arg Gly Leu Arg Ala Ser Val Cys Gln Arg
50 55 60
Leu Ser Met Phe Leu Ala His Phe Leu Arg Gly His Phe Leu Trp Trp
65 70 75 80
Ile Leu Asp Gly Gln Arg Leu Gly Phe Pro Leu Ser Leu Ala Thr Trp
85 90 95
Asn Arg Arg Lys Lys Ser Leu Gln His Leu Leu His Lys His Val Leu
100 105 110
Pro Val Arg Arg His Ala Gly Pro Cys Arg Gly Pro Gln Thr Thr Ala
115 120 125
Arg Gly Pro Arg
130 132

<210> 1197
<211> 64

```
      <212> PRT
      <213> Homo sapiens


      <400> 1197
Met Pro Tyr Leu Ile Leu Phe Phe Ala Val Tyr Ile Leu Tyr Lys Ile
 1               5                  10                  15
Leu Val Lys Val His Leu Phe Ile Ala Glu Ile Ala Leu Tyr Asp Phe
            20                  25                  30
Leu Lys Phe Phe Glu Leu Tyr Gly Ile Cys Met Phe Lys Thr Leu Thr
        35                  40                  45
Cys Leu Val Val Thr Thr Leu Ile Phe Ile Asn Leu Leu Ser Leu  *
    50                  55                  60          63


      <210> 1198
      <211> 53
      <212> PRT
      <213> Homo sapiens


      <400> 1198
Met Leu Gly Pro Pro Glu Ala Arg Leu Ser Leu Cys Ile Leu Leu Trp
 1               5                  10                  15
Ile Ser Ile Leu Cys Pro Trp Tyr Arg Phe Thr Leu Tyr Cys Ser Ser
            20                  25                  30
Trp Pro Tyr Pro Ile Phe Asp Ser Gly Tyr Arg Pro Leu Phe Gly Thr
        35                  40                  45
Thr Leu Leu Phe  *
    50      52


      <210> 1199
      <211> 50
      <212> PRT
      <213> Homo sapiens

      <221> misc_feature
      <222> (1)...(50)
      <223> Xaa = any amino acid or nothing


      <400> 1199
Met Leu Arg Leu Gly Leu Cys Ala Ala Ala Leu Leu Cys Val Cys Arg
 1               5                  10                  15
Pro Gly Ala Val Arg Ala Asp Cys Trp Leu Ile Glu Gly Asp Lys Gly
            20                  25                  30
Tyr Val Trp Leu Ala Ile Cys Asn Gln Asn Gln Pro Ala Tyr Glu Thr
        35                  40                  45
Xaa Pro
    50


      <210> 1200
      <211> 49
      <212> PRT
```

<213> Homo sapiens

<400> 1200
Met Gly Trp Ser Cys Leu Ala Ile Leu Ser Ser Ala Ile Gly His Leu
1 5 10 15
Ile Cys Leu Trp Pro Phe Ala Met Val Val Ala Leu Phe Pro Tyr Leu
20 25 30
Gly Tyr Phe Ser Gly Ser Leu Ser Thr Gln Ile Gly Ser Asp Leu Pro
35 40 45 48
*

<210> 1201
<211> 46
<212> PRT
<213> Homo sapiens

<400> 1201
Met Trp Ala Gly Tyr Val Ile Tyr Thr Leu Phe Cys Arg Phe Ser Phe
1 5 10 15
Ser Leu Ile Ser Ile Arg Ile Arg Lys Leu Gly Ser Ile Gly Phe Glu
20 25 30
Leu Pro Leu Gly Asn Asn Ser Gln Leu Gly Cys Pro Leu *
35 40 45

<210> 1202
<211> 332
<212> PRT
<213> Homo sapiens

<400> 1202
Met Pro Leu Pro Trp Ser Leu Ala Leu Pro Leu Leu Leu Ser Trp Val
1 5 10 15
Ala Gly Gly Phe Gly Asn Ala Ala Ser Ala Arg His His Gly Leu Leu
20 25 30
Ala Ser Ala Arg Gln Pro Gly Val Cys His Tyr Gly Thr Lys Leu Ala
35 40 45
Cys Cys Tyr Gly Trp Arg Arg Asn Ser Lys Gly Val Cys Glu Ala Thr
50 55 60
Cys Glu Pro Gly Cys Lys Phe Gly Glu Cys Val Gly Pro Asn Lys Cys
65 70 75 80
Arg Cys Phe Pro Gly Tyr Thr Gly Lys Thr Cys Ser Gln Asp Val Asn
85 90 95
Glu Cys Gly Met Lys Pro Arg Pro Cys Gln His Arg Cys Val Asn Thr
100 105 110
His Gly Ser Tyr Lys Cys Phe Cys Leu Ser Gly His Met Leu Met Pro
115 120 125
Asp Ala Thr Cys Val Asn Ser Arg Thr Cys Ala Met Ile Asn Cys Gln
130 135 140
Tyr Ser Cys Glu Asp Thr Glu Glu Gly Pro Gln Cys Leu Cys Pro Ser
145 150 155 160
Ser Gly Leu Arg Leu Ala Pro Asn Gly Arg Asp Cys Leu Asp Ile Asp

```
                165                 170                 175
Glu Cys Ala Ser Gly Lys Val Ile Cys Pro Tyr Asn Arg Arg Cys Val
            180                 185                 190
Asn Thr Phe Gly Ser Tyr Tyr Cys Lys Cys His Ile Gly Phe Glu Leu
        195                 200                 205
Gln Tyr Ile Ser Gly Arg Tyr Asp Cys Ile Asp Ile Asn Glu Cys Thr
    210                 215                 220
Met Asp Ser His Thr Cys Ser His His Ala Asn Cys Phe Asn Thr Gln
225                 230                 235                 240
Gly Ser Phe Lys Cys Lys Cys Lys Gln Gly Tyr Lys Gly Asn Gly Leu
                245                 250                 255
Arg Cys Ser Ala Ile Pro Glu Asn Ser Val Lys Glu Val Leu Arg Ala
            260                 265                 270
Pro Gly Thr Ile Lys Asp Arg Ile Lys Lys Leu Leu Ala His Lys Asn
        275                 280                 285
Ser Met Lys Lys Lys Ala Lys Ile Lys Asn Val Thr Pro Glu Pro Thr
    290                 295                 300
Arg Thr Pro Thr Pro Lys Val Asn Leu Gln Pro Phe Asn Tyr Glu Glu
305                 310                 315                 320
Ile Val Ser Arg Gly Gly Asn Ser His Gly Gly  *
                325                 330 331


      <210> 1203
      <211> 825
      <212> PRT
      <213> Homo sapiens


      <400> 1203
Met Ala Arg Leu Gly Asn Cys Ser Leu Thr Trp Ala Ala Leu Ile Ile
 1               5                  10                  15
Leu Leu Leu Pro Gly Ser Leu Glu Glu Cys Gly His Ile Ser Val Ser
            20                  25                  30
Ala Pro Ile Val His Leu Gly Asp Pro Ile Thr Ala Ser Cys Ile Ile
        35                  40                  45
Lys Gln Asn Cys Ser His Leu Asp Pro Glu Pro Gln Ile Leu Trp Arg
    50                  55                  60
Leu Gly Ala Glu Leu Gln Pro Gly Gly Arg Gln Gln Arg Leu Ser Asp
65                  70                  75                  80
Gly Thr Gln Glu Ser Ile Ile Thr Leu Pro His Leu Asn His Thr Gln
                85                  90                  95
Ala Phe Leu Ser Cys Cys Leu Asn Trp Gly Asn Ser Leu Gln Ile Leu
            100                 105                 110
Asp Gln Val Glu Leu Arg Ala Gly Tyr Pro Pro Ala Ile Pro His Asn
        115                 120                 125
Leu Ser Cys Leu Met Asn Leu Thr Thr Ser Ser Leu Ile Cys Gln Trp
    130                 135                 140
Glu Pro Gly Pro Glu Thr His Leu Pro Thr Ser Phe Thr Leu Lys Ser
145                 150                 155                 160
Phe Lys Ser Arg Gly Asn Cys Gln Thr Gln Gly Asp Ser Ile Leu Asp
                165                 170                 175
Cys Val Pro Lys Asp Gly Gln Ser His Cys Cys Ile Pro Arg Lys His
            180                 185                 190
Leu Leu Leu Tyr Gln Asn Met Gly Ile Trp Val Gln Ala Glu Asn Ala
        195                 200                 205
Leu Gly Thr Ser Met Ser Pro Gln Leu Cys Leu Asp Pro Met Asp Val
    210                 215                 220
```

Val Lys Leu Glu Pro Pro Met Leu Arg Thr Met Asp Pro Ser Pro Glu
225 230 235 240
Ala Ala Pro Pro Gln Ala Gly Cys Leu Gln Leu Cys Trp Glu Pro Trp
245 250 255
Gln Pro Gly Leu His Ile Asn Gln Lys Cys Glu Leu Arg His Lys Pro
260 265 270
Gln Arg Gly Glu Ala Ser Trp Ala Leu Val Gly Pro Leu Pro Leu Glu
275 280 285
Ala Leu Gln Tyr Glu Leu Cys Gly Leu Leu Pro Ala Thr Ala Tyr Thr
290 295 300
Leu Gln Ile Arg Cys Ile Arg Trp Pro Leu Pro Gly His Trp Ser Asp
305 310 315 320
Trp Ser Pro Ser Leu Glu Leu Arg Thr Thr Glu Arg Ala Pro Thr Val
325 330 335
Arg Leu Asp Thr Trp Trp Arg Gln Arg Gln Leu Asp Pro Arg Thr Val
340 345 350
Gln Leu Phe Trp Lys Pro Val Pro Leu Glu Glu Asp Ser Gly Arg Ile
355 360 365
Gln Gly Tyr Val Val Ser Trp Arg Pro Ser Gly Gln Ala Gly Ala Ile
370 375 380
Leu Pro Leu Cys Asn Thr Thr Glu Leu Ser Cys Thr Phe His Leu Pro
385 390 395 400
Ser Glu Ala Gln Glu Val Ala Leu Val Ala Tyr Asn Ser Ala Gly Thr
405 410 415
Ser Arg Pro Thr Pro Val Val Phe Ser Glu Ser Arg Gly Pro Ala Leu
420 425 430
Thr Arg Leu His Ala Met Ala Arg Asp Pro His Ser Leu Trp Val Gly
435 440 445
Trp Glu Pro Pro Asn Pro Trp Pro Gln Gly Tyr Val Ile Glu Trp Gly
450 455 460
Leu Gly Pro Pro Ser Ala Ser Asn Ser Asn Lys Thr Trp Arg Met Glu
465 470 475 480
Gln Asn Gly Arg Ala Thr Gly Phe Leu Leu Lys Glu Asn Ile Arg Pro
485 490 495
Phe Gln Leu Tyr Glu Ile Ile Val Thr Pro Leu Tyr Gln Asp Thr Met
500 505 510
Gly Pro Ser Gln His Val Tyr Ala Tyr Ser Gln Glu Met Ala Pro Ser
515 520 525
His Ala Pro Glu Leu His Leu Lys His Ile Gly Lys Thr Trp Ala Gln
530 535 540
Leu Glu Trp Val Pro Glu Pro Pro Glu Leu Gly Lys Ser Pro Leu Thr
545 550 555 560
His Tyr Thr Ile Phe Trp Thr Asn Ala Gln Asn Gln Ser Phe Ser Ala
565 570 575
Ile Leu Asn Ala Ser Ser Arg Gly Phe Val Leu His Gly Leu Glu Pro
580 585 590
Ala Ser Leu Tyr His Ile His Leu Met Ala Ala Ser Gln Ala Gly Ala
595 600 605
Thr Asn Ser Thr Val Leu Thr Leu Met Thr Leu Thr Pro Ala Pro Thr
610 615 620
Gly Arg Ile Pro Ser Gly Gln Val Ser Gln Thr Gln Leu Thr Ala Ala
625 630 635 640
Trp Ala Pro Gly Cys Pro Gln Ser Trp Arg Arg Met Pro Ser Ser Cys
645 650 655
Pro Ala Leu Ala Arg His Pro Ser Pro Ser Ser Gln Cys Trp Arg Arg
660 665 670
Met Lys Arg Ser Arg Cys Pro Gly Ser Pro Ile Thr Ala Gln Arg Pro
675 680 685
Val Ala Ser Pro Leu Trp Ser Arg Pro Met Cys Ser Arg Gly Thr Gln

```
    690                 695                 700
Glu Gln Phe Pro Pro Ser Pro Asn Pro Ser Leu Ala Pro Ala Ile Arg
705                 710                 715                 720
Ser Phe Met Gly Ser Cys Trp Ala Ala Pro Gln Ala Gln Gly Gln Gly
                725                 730                 735
Thr Ile Ser Ala Val Thr Pro Leu Ser Pro Ser Trp Arg Ala Ser Pro
            740                 745                 750
Pro Ala Pro Ser Pro Met Arg Thr Ser Gly Ser Arg Pro Ala Pro Trp
        755                 760                 765
Gly Pro Leu Val Thr Pro Ser Pro Lys Ser Gln Glu Asp Asp Cys Val
    770                 775                 780
Phe Gly Pro Leu Leu Asn Phe Pro Pro Ser Cys Arg Gly Ser Gly Ser
785                 790                 795                 800
Met Gly Trp Arg Arg Trp Gly Ala Ser Arg Ala Ser Leu Gly Phe Pro
                805                 810                 815
Ser Trp Ala Cys Leu Leu Lys Ala  *
            820                 824
```

```
     <210> 1204
     <211> 48
     <212> PRT
     <213> Homo sapiens


     <400> 1204
Met Leu Leu Phe Ser Ser Arg Phe Ile Met Phe Leu Trp Pro Pro Val
 1               5                  10                  15
Ser Gly Val Cys Leu Ser Phe Ile Arg Asp Arg Ser Phe Leu Pro Met
            20                  25                  30
Cys His Phe Ile Tyr Val Leu Ile Leu Cys Asn Ser Ile Ala Leu  *
        35                  40                  45      47
```

```
     <210> 1205
     <211> 46
     <212> PRT
     <213> Homo sapiens


     <400> 1205
Met Gly Ser Phe Ser Phe Ile Leu Val Leu Phe Ile Asp Cys Leu Cys
 1               5                  10                  15
Met Phe Pro Ser Val Leu Val Gln Leu Leu Cys Thr Tyr Ser Ser Leu
            20                  25                  30
Met Lys Thr Pro Leu Trp Leu Gln Ala Arg Ser Ser His  *
        35                  40                  45
```

```
     <210> 1206
     <211> 88
     <212> PRT
     <213> Homo sapiens


     <400> 1206
```

Met Gln Trp Cys Asn Leu Thr Ala Thr Ser Ala Phe Gln Ile Glu Ala
1 5 10 15
Ile Leu Leu Pro Gln Leu Ser Pro Val Ala Gly Ile Thr Gly Thr Cys
20 25 30
Tyr His Ala Trp Leu Ile Phe Val Phe Leu Val Glu Thr Gly Phe His
35 40 45
His Val Gly Gln Ala Gly Leu Glu Leu Leu Thr Ser Gly Asp Pro Pro
50 55 60
Thr Leu Ala Ser Gln Ser Ala Gly Ile Thr Ser Val Ser His His Ala
65 70 75 80
Gln Pro Leu Lys Gly Thr Phe *
85 87

<210> 1207
<211> 186
<212> PRT
<213> Homo sapiens

<400> 1207
Met Ile Leu Asn Lys Ala Leu Met Leu Gly Ala Leu Ala Leu Thr Thr
1 5 10 15
Val Met Ser Pro Cys Gly Gly Glu Asp Ile Val Ala Asp His Val Ala
20 25 30
Ser Tyr Gly Val Asn Leu Tyr Gln Ser Tyr Gly Pro Ser Gly Gln Tyr
35 40 45
Ser His Glu Phe Asp Gly Asp Glu Glu Phe Tyr Val Asp Leu Glu Arg
50 55 60
Lys Glu Thr Val Trp Gln Leu Pro Leu Phe Arg Arg Phe Arg Arg Phe
65 70 75 80
Asp Pro Gln Phe Ala Leu Thr Asn Ile Ala Val Leu Lys His Asn Leu
85 90 95
Asn Ile Val Ile Lys Arg Ser Asn Ser Thr Ala Ala Thr Asn Glu Val
100 105 110
Pro Glu Val Thr Val Phe Ser Lys Ser Pro Val Thr Leu Gly Gln Pro
115 120 125
Asn Thr Leu Ile Cys Leu Val Asp Asn Ile Phe Pro Pro Val Val Asn
130 135 140
Ile Thr Trp Leu Ser Asn Gly His Ser Val Thr Glu Gly Val Ser Glu
145 150 155 160
Thr Arg Pro Ser Ser Pro Lys Ser Asp His Phe Leu Leu Gln Asp Gln
165 170 175
Val Thr Ser Pro Ser Phe Pro Phe Glu *
180 185

<210> 1208
<211> 46
<212> PRT
<213> Homo sapiens

<400> 1208
Met Asn Pro His Leu Gly Val Phe Leu Val Leu Val Ser Phe Phe Leu
1 5 10 15
Ser Leu Leu Asp Ser Gln Leu His Ser Trp Ile Val Leu His Asn Ser

```
            20                      25                  30
Pro Ser Ser Arg Met Trp Lys Ser Ile Ile Phe Phe Leu  *
        35                  40                  45
```

<210> 1209
<211> 199
<212> PRT
<213> Homo sapiens

<400> 1209

```
Met Ala Leu Leu Val Pro Leu Ala Leu Leu Val Ile Gln Ala His Leu
 1               5                  10                  15
Val Leu Ser Val Gln Leu Glu Arg Val Val Thr Glu Glu Lys Val Ala
            20                  25                  30
Leu Leu Ala Leu Leu Val Leu Pro Val Leu Leu Val Pro Glu Val Leu
        35                  40                  45
Leu Val Leu Lys Ala His Val Val Thr Lys Val Lys Gln Val Asn Val
    50                  55                  60
Glu Leu Leu Ala Ser Lys Asp Ile Glu Asp Ser Leu Val Ile Gln Val
65                  70                  75                  80
Pro Gln Val Leu Gln Ala Leu Leu Val Ser Arg Val Gln Ser Ala Val
                85                  90                  95
Gln Asp Leu Gln Ala Pro Glu Asp Leu Leu Asp Pro Val Asp Leu Leu
            100                 105                 110
Ala Lys Met Glu Pro Val Asp Ile Gln Val Pro Leu Asp His Gln Gly
        115                 120                 125
Leu Glu Val Thr Glu Val Lys Glu Asp Leu Arg Ala Pro Gln Ala Thr
    130                 135                 140
Gln Gly Asn Gln Ala Leu Leu Asp Leu Leu Val Pro Leu Val Leu Ala
145                 150                 155                 160
Val Val Val Leu Glu Pro Leu Pro Leu Leu Gly Leu Glu Val Lys Lys
                165                 170                 175
Leu Ala Val Leu Pro Arg Ile Met Glu Met Asn Gln Trp Ile Ser Lys
            180                 185                 190
Ser Thr Pro Met Arg Leu  *
        195         198
```

<210> 1210
<211> 59
<212> PRT
<213> Homo sapiens

<400> 1210

```
Met Leu Val Thr Arg Pro Ser Gly Asn Thr Trp Ile Pro Phe Phe Cys
 1               5                  10                  15
Trp Leu Leu Phe Cys Val Val Glu Leu Leu Ser Pro Gly Asn Leu Gly
            20                  25                  30
Pro Ser Val Leu Glu Val Val Leu Pro Asp Val Phe Lys Leu Asp Leu
        35                  40                  45
Leu Ser Ser Leu Leu Asp Val Gly Ser Leu  *
    50                  55          58
```

<210> 1211
<211> 227
<212> PRT
<213> Homo sapiens

<221> misc_feature
<222> (1)...(227)
<223> Xaa = any amino acid or nothing

<400> 1211
Met Ala Ser Ile Cys Ser Trp Arg Val Met Leu Ala Trp Ala Ala Cys
1 5 10 15
Trp Val Arg Ala His Ala Ala Leu Ser Gly His Pro Arg Ser Thr Phe
20 25 30
Ser Leu Trp Leu Ser Gly Ile Ser Leu Pro Xaa Pro Ile Phe Leu Pro
35 40 45
Met Ala Val Ser Leu Leu Thr Pro Lys Asp Val Lys Tyr Ala Arg Ser
50 55 60
Pro Asn Cys Phe Lys Ala Ala Leu Asn Ile Pro Asp Pro Gly Ala Val
65 70 75 80
His Leu Ile Ile Ala Leu Leu Leu Thr Asp Gly Ala Ile Pro Leu Leu
85 90 95
Gln Pro Ala Arg Val Lys Lys Ser Asn Ala His Val Phe Leu His Phe
100 105 110
Ala Gly Gly Asp Leu Leu Pro Ser Asn Gly Gly His Lys Ile Leu Ile
115 120 125
Trp Ser Arg Gly Trp Arg Gln Gly Leu Gly Gly Phe Gly Ile Ile Ile
130 135 140
Leu Ala Asp Asn Asp Leu Val Trp Ser Trp Gly Gln Ser Trp Arg His
145 150 155 160
Gly Cys Leu Leu Gly Val Gly Ala Leu Ser Ala Leu Leu Leu His His
165 170 175
Leu Asn Pro His Pro Tyr Leu Val Leu Gly Cys Pro Gly Pro Ala Gly
180 185 190
Lys Glu Ala Pro Pro Pro Ser Pro Val Cys His Pro Pro His Gln Thr
195 200 205
Arg Pro Pro Ser Gln Leu Pro His Ser Pro Gln Thr Phe His Ser Ala
210 215 220
Pro Glu *
225 226

<210> 1212
<211> 62
<212> PRT
<213> Homo sapiens

<400> 1212
Met Cys Val Ser Val Arg Val Cys Val Cys Val Cys Val Cys Ala Arg
1 5 10 15
Val Cys Ala Arg Leu Cys Val Cys Val His Ala Arg Leu Cys Val His
20 25 30
Val Arg Val Ser Ala Arg Val Ser Val Tyr Val Cys Thr Arg Val Ser
35 40 45
Val Cys Val His Ala Arg Ala Arg His His Arg Ser Ile *

```
     50                  55                  60  61


     <210> 1213
     <211> 55
     <212> PRT
     <213> Homo sapiens


     <400> 1213
Met Phe Arg Arg Leu Thr Phe Ala Gln Leu Leu Phe Ala Thr Val Leu
  1               5                  10                  15
Gly Ile Ala Gly Gly Val Tyr Ile Phe Gln Pro Val Phe Glu Gln Tyr
             20                  25                  30
Ala Lys Asp Gln Lys Glu Leu Lys Glu Lys Met Gln Leu Val Gln Glu
         35                  40                  45
Ser Glu Glu Lys Lys Ser  *
     50              54


     <210> 1214
     <211> 642
     <212> PRT
     <213> Homo sapiens


     <400> 1214
Met Thr Met Tyr Leu Trp Leu Lys Leu Leu Ala Phe Gly Phe Ala Phe
  1               5                  10                  15
Leu Asp Thr Glu Val Phe Val Thr Gly Gln Ser Pro Thr Pro Ser Pro
             20                  25                  30
Thr Asp Ala Tyr Leu Asn Ala Ser Glu Thr Thr Thr Leu Ser Pro Ser
         35                  40                  45
Gly Ser Ala Val Ile Ser Thr Thr Thr Ile Ala Thr Thr Pro Ser Lys
     50                  55                  60
Pro Thr Cys Asp Glu Lys Tyr Ala Asn Ile Thr Val Asp Tyr Leu Tyr
 65                  70                  75                  80
Asn Lys Glu Thr Lys Leu Phe Thr Ala Lys Leu Asn Val Asn Glu Asn
                 85                  90                  95
Val Glu Cys Gly Asn Asn Thr Cys Thr Asn Asn Glu Val His Asn Leu
            100                 105                 110
Thr Glu Cys Lys Asn Ala Ser Val Ser Ile Ser His Asn Ser Cys Thr
        115                 120                 125
Ala Pro Asp Lys Thr Leu Ile Leu Asp Val Pro Pro Gly Val Glu Lys
    130                 135                 140
Phe Gln Leu His Asp Cys Thr Gln Val Glu Lys Ala Asp Thr Thr Ile
145                 150                 155                 160
Cys Leu Lys Trp Lys Asn Ile Glu Thr Phe Thr Cys Asp Thr Gln Asn
                165                 170                 175
Ile Thr Tyr Arg Phe Gln Cys Gly Asn Met Ile Phe Asp Asn Lys Glu
            180                 185                 190
Ile Lys Leu Glu Asn Leu Glu Pro Glu His Glu Tyr Lys Cys Asp Ser
        195                 200                 205
Glu Ile Leu Tyr Asn Asn His Lys Phe Thr Asn Ala Ser Lys Ile Ile
    210                 215                 220
Lys Thr Asp Phe Gly Ser Pro Gly Glu Pro Gln Ile Ile Phe Cys Arg
225                 230                 235                 240
```

```
Ser Glu Ala Ala His Gln Gly Val Ile Thr Trp Asn Pro Pro Gln Arg
                245                 250                 255
Ser Phe His Asn Phe Thr Leu Cys Tyr Ile Lys Glu Thr Glu Lys Asp
            260                 265                 270
Cys Leu Asn Leu Asp Lys Asn Leu Ile Lys Tyr Asp Leu Gln Asn Leu
        275                 280                 285
Lys Pro Tyr Thr Lys Tyr Val Leu Ser Leu His Ala Tyr Ile Ile Ala
    290                 295                 300
Lys Val Gln Arg Asn Gly Ser Ala Ala Met Cys His Phe Thr Thr Lys
305                 310                 315                 320
Ser Ala Pro Pro Ser Gln Val Trp Asn Met Thr Val Ser Met Thr Ser
                325                 330                 335
Asp Asn Ser Met His Val Lys Cys Arg Pro Pro Arg Asp Arg Asn Gly
            340                 345                 350
Pro His Glu Arg Tyr His Leu Glu Val Glu Ala Gly Asn Thr Leu Val
        355                 360                 365
Arg Asn Glu Ser His Lys Asn Cys Asp Phe Arg Val Lys Asp Leu Gln
    370                 375                 380
Tyr Ser Thr Asp Tyr Thr Phe Lys Ala Tyr Phe His Asn Gly Asp Tyr
385                 390                 395                 400
Pro Gly Glu Pro Phe Ile Leu His His Ser Thr Ser Tyr Asn Ser Lys
                405                 410                 415
Ala Leu Ile Ala Phe Leu Ala Phe Leu Ile Ile Val Thr Ser Ile Ala
            420                 425                 430
Leu Leu Val Val Leu Tyr Lys Ile Tyr Asp Leu His Lys Lys Arg Ser
        435                 440                 445
Cys Asn Leu Asp Glu Gln Gln Glu Leu Val Glu Arg Asp Asp Glu Lys
    450                 455                 460
Gln Leu Met Asn Val Glu Pro Ile His Ala Asp Ile Leu Leu Glu Thr
465                 470                 475                 480
Tyr Lys Arg Lys Ile Ala Asp Glu Gly Arg Leu Phe Leu Ala Glu Phe
                485                 490                 495
Gln Ser Ile Pro Arg Val Phe Ser Lys Phe Pro Ile Lys Glu Ala Arg
            500                 505                 510
Lys Pro Phe Asn Gln Asn Lys Asn Arg Tyr Val Asp Ile Leu Pro Tyr
        515                 520                 525
Asp Tyr Asn Arg Val Glu Leu Ser Glu Ile Asn Gly Asp Ala Gly Ser
    530                 535                 540
Asn Tyr Ile Asn Ala Ser Tyr Ile Asp Gly Phe Lys Glu Pro Arg Lys
545                 550                 555                 560
Tyr Ile Ala Ala Gln Gly Pro Arg Asp Glu Thr Val Asp Asp Phe Trp
                565                 570                 575
Arg Met Ile Trp Glu Gln Lys Ala Thr Val Ile Val Met Val Thr Arg
            580                 585                 590
Cys Glu Glu Gly Asn Arg Asn Lys Cys Ala Glu Tyr Trp Pro Ser Met
        595                 600                 605
Glu Glu Gly Thr Arg Ala Phe Gly Glu Cys Cys Cys Lys Asp Leu Thr
    610                 615                 620
Lys His Lys Arg Cys Pro Arg Leu His His Ser Glu Ile Glu His Cys
625                 630                 635                 640
Lys  *
641
```

<210> 1215
<211> 85
<212> PRT
<213> Homo sapiens

<400> 1215
Met Leu Phe Leu Thr Leu Ile Ser Phe Cys Gly Phe Leu Leu Leu His
1 5 10 15
Arg Leu Thr Ser Met Val Arg Leu Phe Leu Gly Ala Ala Ile Gln Lys
20 25 30
Ile Leu Ser Lys Arg Leu Glu Phe Ser Leu Leu Pro Leu Val Ser Phe
35 40 45
Ala Gly Ser Val Asn Met Ala Gly Pro Cys Thr Ala Asn Ala Gly Pro
50 55 60
His Gly Gly Leu Gly Lys Pro Gly Arg Leu Cys Gly Ser Phe Arg Ser
65 70 75 80
Ser Arg Ser Gln *
84

<210> 1216
<211> 403
<212> PRT
<213> Homo sapiens

<400> 1216
Met Ala Ser Val Val Leu Pro Ser Gly Ser Gln Cys Ala Ala Ala Ala
1 5 10 15
Ala Ala Ala Ala Pro Pro Gly Leu Arg Leu Arg Leu Leu Leu Leu Leu
20 25 30
Phe Ser Ala Ala Ala Leu Ile Pro Thr Gly Asp Gly Gln Asn Leu Phe
35 40 45
Thr Lys Asp Val Thr Val Ile Glu Gly Glu Val Ala Thr Ile Ser Cys
50 55 60
Gln Val Asn Lys Ser Asp Asp Ser Val Ile Gln Leu Leu Asn Pro Asn
65 70 75 80
Arg Gln Thr Ile Tyr Phe Arg Asp Phe Arg Pro Leu Lys Asp Ser Arg
85 90 95
Phe Gln Leu Leu Asn Phe Ser Ser Ser Glu Leu Lys Val Ser Leu Thr
100 105 110
Asn Val Ser Ile Ser Asp Glu Gly Arg Tyr Phe Cys Gln Leu Tyr Thr
115 120 125
Asp Pro Pro Gln Glu Ser Tyr Thr Thr Ile Thr Val Leu Val Pro Pro
130 135 140
Arg Asn Leu Met Ile Asp Ile Gln Lys Asp Thr Ala Val Glu Gly Glu
145 150 155 160
Glu Ile Glu Val Asn Cys Thr Ala Met Ala Ser Lys Pro Ala Thr Thr
165 170 175
Ile Arg Trp Phe Lys Gly Asn Thr Glu Leu Lys Gly Lys Ser Glu Val
180 185 190
Glu Glu Trp Ser Asp Met Tyr Thr Val Thr Ser Gln Leu Met Leu Lys
195 200 205
Val His Lys Glu Asp Asp Gly Val Pro Val Ile Cys Gln Val Glu His
210 215 220
Pro Ala Val Thr Gly Asn Leu Gln Thr Gln Arg Tyr Leu Glu Val Gln
225 230 235 240
Tyr Lys Pro Gln Val His Ile Gln Met Thr Tyr Pro Leu Gln Gly Leu
245 250 255
Thr Arg Glu Gly Asp Ala Leu Glu Leu Thr Cys Glu Ala Ile Gly Lys
260 265 270

```
Pro Gln Pro Val Met Val Thr Trp Val Arg Val Asp Asp Glu Met Pro
        275                 280                 285
Gln His Ala Val Leu Ser Gly Pro Asn Leu Phe Ile Asn Asn Leu Asn
    290                 295                 300
Lys Thr Asp Asn Gly Thr Tyr Arg Cys Glu Ala Ser Asn Ile Val Gly
305                 310                 315                 320
Lys Ala His Ser Asp Tyr Met Leu Tyr Val Tyr Asp Pro Pro Thr Thr
                325                 330                 335
Ile Pro Pro Pro Thr Thr Thr Thr Thr Thr Thr Thr Thr Thr Thr Thr
            340                 345                 350
Thr Ile Leu Thr Ile Ile Thr Asp Ser Arg Ala Gly Glu Glu Gly Ser
        355                 360                 365
Ile Arg Ala Val Asp His Ala Val Ile Gly Gly Val Val Ala Val Val
    370                 375                 380
Val Phe Ala Met Leu Cys Leu Leu Ile Ile Leu Gly Arg Tyr Phe Ala
385                 390                 395                 400
Gln Thr  *
    402
```

```
<210> 1217
<211> 49
<212> PRT
<213> Homo sapiens
```

```
<400> 1217
Met Arg Ala Trp Ala Trp Pro Phe Cys Thr Ser Val Thr Ser Leu Ser
 1               5                  10                  15
Ala Met Ala Ser Pro Trp Arg Arg Trp Pro Arg Arg Pro Ala Ser Arg
            20                  25                  30
Thr Ala Ser Arg Ala Pro Ser Ala Gly Ile Ser Gly Ser Thr Ala Pro
        35                  40                  45              48
 *
```

```
<210> 1218
<211> 304
<212> PRT
<213> Homo sapiens
```

```
<400> 1218
Met Ala Arg Arg Ser Arg His Arg Leu Leu Leu Leu Leu Leu Arg Tyr
 1               5                  10                  15
Leu Val Val Ala Leu Gly Tyr His Lys Ala Tyr Gly Phe Ser Ala Pro
            20                  25                  30
Lys Asp Gln Gln Val Val Thr Ala Val Glu Tyr Gln Glu Ala Ile Leu
        35                  40                  45
Ala Cys Lys Thr Pro Lys Lys Thr Val Ser Ser Arg Leu Glu Trp Lys
    50                  55                  60
Lys Leu Gly Arg Ser Val Ser Phe Val Tyr Tyr Gln Gln Thr Leu Gln
65                  70                  75                  80
Gly Asp Phe Lys Asn Arg Ala Glu Met Ile Asp Phe Asn Ile Arg Ile
                85                  90                  95
Lys Asn Val Thr Arg Ser Asp Ala Gly Lys Tyr Arg Cys Glu Val Ser
```

```
            100                 105                 110
Ala Pro Ser Glu Gln Gly Gln Asn Leu Glu Glu Asp Thr Val Thr Leu
        115                 120                 125
Glu Val Leu Gly Asp Val His Val Leu Ala Pro Ala Val Pro Ser Cys
    130                 135                 140
Glu Val Pro Ser Ser Ala Leu Ser Gly Thr Val Val Glu Leu Arg Cys
145                 150                 155                 160
Gln Asp Lys Glu Gly Asn Pro Ala Pro Glu Tyr Thr Trp Phe Lys Asp
                165                 170                 175
Gly Ile Arg Leu Leu Glu Asn Pro Arg Leu Gly Ser Gln Ser Thr Asn
            180                 185                 190
Ser Ser Tyr Thr Met Asn Thr Lys Thr Gly Thr Leu Gln Phe Asn Thr
        195                 200                 205
Val Ser Lys Leu Asp Thr Gly Glu Tyr Ser Cys Glu Ala Arg Asn Ser
    210                 215                 220
Val Gly Tyr Arg Arg Cys Pro Gly Lys Arg Met Gln Val Asp Asp Leu
225                 230                 235                 240
Asn Ile Ser Gly Ile Ile Ala Ala Val Val Val Val Ala Leu Val Ile
                245                 250                 255
Ser Val Cys Gly Leu Gly Val Cys Tyr Ala Gln Arg Lys Gly Tyr Phe
            260                 265                 270
Ser Lys Glu Thr Ser Phe Gln Lys Ser Asn Ser Ser Ser Lys Ala Thr
        275                 280                 285
Thr Met Ser Glu Asn Asp Phe Lys His Thr Lys Ser Phe Ile Ile  *
    290                 295                 300         303


     <210> 1219
     <211> 1126
     <212> PRT
     <213> Homo sapiens


     <400> 1219
Met Trp Phe Leu Phe Leu Cys Pro Asn Leu Trp Ala Met Pro Val Gln
  1               5                  10                  15
Ile Ile Met Gly Val Ile Leu Leu Tyr Asn Leu Leu Gly Ser Ser Ala
            20                  25                  30
Leu Val Gly Ala Ala Val Ile Val Leu Leu Ala Pro Ile Gln Tyr Phe
        35                  40                  45
Ile Ala Thr Lys Leu Ala Glu Ala Gln Lys Ser Thr Leu Asp Tyr Ser
    50                  55                  60
Thr Glu Arg Leu Lys Lys Thr Asn Glu Ile Leu Lys Gly Ile Lys Leu
 65                  70                  75                  80
Leu Lys Leu Tyr Ala Trp Glu His Ile Phe Cys Lys Ser Val Glu Glu
                85                  90                  95
Thr Arg Met Lys Glu Leu Ser Ser Leu Lys Thr Phe Ala Leu Tyr Thr
            100                 105                 110
Ser Leu Ser Ile Phe Met Asn Ala Ala Ile Pro Ile Ala Ala Val Leu
        115                 120                 125
Ala Thr Phe Val Thr His Ala Tyr Ala Ser Gly Asn Asn Leu Lys Pro
    130                 135                 140
Ala Glu Ala Phe Ala Ser Leu Ser Leu Phe His Ile Leu Val Thr Pro
145                 150                 155                 160
Leu Phe Leu Leu Ser Thr Val Val Arg Phe Ala Val Lys Ala Ile Ile
                165                 170                 175
Ser Val Gln Lys Leu Asn Glu Phe Leu Leu Ser Asp Glu Ile Gly Asp
            180                 185                 190
```

Asp Ser Trp Arg Thr Gly Glu Ser Ser Leu Pro Phe Glu Ser Cys Lys
195 200 205
Lys His Thr Gly Val Gln Pro Lys Thr Ile Asn Arg Lys Gln Pro Gly
210 215 220
Arg Tyr His Leu Asp Ser Tyr Glu Gln Ser Thr Arg Arg Leu Arg Pro
225 230 235 240
Ala Glu Thr Glu Asp Ile Ala Ile Lys Val Thr Asn Gly Tyr Phe Ser
245 250 255
Trp Gly Ser Gly Leu Ala Thr Leu Ser Asn Ile Asp Ile Arg Ile Pro
260 265 270
Thr Gly Gln Leu Thr Met Ile Val Gly Gln Val Gly Cys Gly Lys Ser
275 280 285
Ser Leu Leu Leu Ala Ile Leu Gly Glu Met Gln Thr Leu Glu Gly Lys
290 295 300
Val His Trp Ser Asn Val Asn Glu Ser Glu Pro Ser Phe Glu Ala Thr
305 310 315 320
Arg Ser Arg Asn Arg Tyr Ser Val Ala Tyr Ala Ala Gln Lys Pro Trp
325 330 335
Leu Leu Asn Ala Thr Val Glu Glu Asn Ile Thr Phe Gly Ser Pro Phe
340 345 350
Asn Lys Gln Arg Tyr Lys Ala Val Thr Asp Ala Cys Ser Leu Gln Pro
355 360 365
Asp Ile Asp Leu Leu Pro Phe Gly Asp Gln Thr Glu Ile Gly Glu Arg
370 375 380
Gly Ile Asn Leu Ser Gly Gly Gln Arg Gln Arg Ile Cys Val Ala Arg
385 390 395 400
Ala Leu Tyr Gln Asn Thr Asn Ile Val Phe Leu Asp Asp Pro Phe Ser
405 410 415
Ala Leu Asp Ile His Leu Ser Asp His Leu Met Gln Glu Gly Ile Leu
420 425 430
Lys Phe Leu Gln Asp Asp Lys Arg Thr Leu Val Leu Val Thr His Lys
435 440 445
Leu Gln Tyr Leu Thr His Ala Asp Trp Ile Ile Ala Met Lys Asp Gly
450 455 460
Ser Val Leu Arg Glu Gly Thr Leu Lys Asp Ile Gln Thr Lys Asp Val
465 470 475 480
Glu Leu Tyr Glu His Trp Lys Thr Leu Met Asn Arg Gln Asp Gln Glu
485 490 495
Leu Glu Lys Asp Met Glu Ala Asp Gln Thr Thr Leu Glu Arg Lys Thr
500 505 510
Leu Arg Arg Ala Met Tyr Ser Arg Glu Ala Lys Ala Gln Met Glu Asp
515 520 525
Glu Asp Glu Glu Glu Glu Glu Glu Glu Asp Glu Asp Asp Asn Met Ser
530 535 540
Thr Val Met Arg Leu Arg Thr Lys Met Pro Trp Lys Thr Cys Trp Arg
545 550 555 560
Tyr Leu Thr Ser Gly Gly Phe Phe Leu Leu Ile Leu Met Ile Phe Ser
565 570 575
Lys Leu Leu Lys His Ser Val Ile Val Ala Ile Asp Tyr Trp Leu Ala
580 585 590
Thr Trp Thr Ser Glu Tyr Ser Ile Asn Asn Thr Gly Lys Ala Asp Gln
595 600 605
Thr Tyr Tyr Val Ala Gly Phe Ser Ile Leu Cys Gly Ala Gly Ile Phe
610 615 620
Leu Cys Leu Val Thr Ser Leu Thr Val Glu Trp Met Gly Leu Thr Ala
625 630 635 640
Ala Lys Asn Leu His His Asn Leu Leu Asn Lys Ile Ile Leu Gly Pro
645 650 655
Ile Arg Phe Phe Asp Thr Thr Pro Leu Gly Leu Ile Leu Asn Arg Phe

```
            660                     665                     670
Ser Ala Asp Thr Asn Ile Ile Asp Gln His Ile Pro Pro Thr Leu Glu
        675                     680                     685
Ser Leu Thr Arg Ser Thr Leu Leu Cys Leu Ser Ala Ile Gly Met Ile
    690                     695                     700
Ser Tyr Ala Thr Pro Val Phe Leu Val Ala Leu Leu Pro Leu Gly Val
705                     710                     715                     720
Ala Phe Tyr Phe Ile Gln Lys Tyr Phe Arg Val Ala Ser Lys Asp Leu
                725                     730                     735
Gln Glu Leu Asp Asp Ser Thr Gln Leu Pro Leu Leu Cys His Phe Ser
            740                     745                     750
Glu Thr Ala Glu Gly Leu Thr Thr Ile Arg Ala Phe Arg His Glu Thr
        755                     760                     765
Arg Phe Lys Gln Arg Met Leu Glu Leu Thr Asp Thr Asn Asn Ile Ala
    770                     775                     780
Tyr Leu Phe Leu Ser Ala Ala Asn Arg Trp Leu Glu Val Arg Thr Asp
785                     790                     795                     800
Tyr Leu Gly Ala Cys Ile Val Leu Thr Ala Ser Ile Ala Ser Ile Ser
                805                     810                     815
Gly Ser Ser Asn Ser Gly Leu Val Gly Leu Gly Leu Leu Tyr Ala Leu
            820                     825                     830
Thr Ile Thr Asn Tyr Leu Asn Trp Val Val Arg Asn Leu Ala Asp Leu
        835                     840                     845
Glu Val Gln Met Gly Ala Val Lys Lys Val Asn Ser Phe Leu Thr Met
    850                     855                     860
Glu Ser Glu Asn Tyr Glu Gly Thr Met Asp Pro Ser Gln Val Pro Glu
865                     870                     875                     880
His Trp Pro Gln Glu Gly Glu Ile Lys Ile His Asp Leu Cys Val Arg
                885                     890                     895
Tyr Glu Asn Asn Leu Lys Pro Val Leu Lys His Val Lys Ala Tyr Ile
            900                     905                     910
Lys Pro Gly Gln Lys Val Gly Ile Cys Gly Arg Thr Gly Ser Gly Lys
        915                     920                     925
Ser Ser Leu Ser Leu Ala Phe Phe Arg Met Val Asp Ile Phe Asp Gly
    930                     935                     940
Lys Ile Val Ile Asp Gly Ile Asp Ile Ser Lys Leu Pro Leu His Thr
945                     950                     955                     960
Leu Arg Ser Arg Leu Ser Ile Ile Leu Gln Asp Pro Ile Leu Phe Ser
                965                     970                     975
Gly Ser Ile Arg Phe Asn Leu Asp Pro Glu Cys Lys Cys Thr Asp Asp
            980                     985                     990
Arg Leu Trp Glu Ala Leu Glu Ile Ala Gln Leu Lys Asn Met Val Lys
        995                     1000                    1005
Ser Leu Pro Gly Gly Leu Asp Ala Val Val Thr Glu Gly Gly Glu Asn
    1010                    1015                    1020
Phe Ser Val Gly Gln Arg Gln Leu Phe Cys Leu Ala Arg Ala Phe Val
1025                    1030                    1035                    1040
Arg Lys Ser Ser Ile Leu Ile Met Asp Glu Ala Thr Ala Ser Ile Asp
                1045                    1050                    1055
Met Ala Thr Glu Asn Ile Leu Gln Lys Val Val Met Thr Ala Phe Ala
            1060                    1065                    1070
Asp Arg Thr Val Val Thr Met Ala His Arg Val Ser Ser Ile Met Asp
        1075                    1080                    1085
Ala Gly Leu Val Leu Val Phe Ser Glu Gly Ile Leu Val Glu Cys Asp
    1090                    1095                    1100
Thr Val Pro Asn Leu Phe Ala His Lys Asn Gly Pro Phe Ser Thr Leu
1105                    1110                    1115                    1120
Val Met Thr Asn Lys  *
                1125
```

<210> 1220
<211> 46
<212> PRT
<213> Homo sapiens

<400> 1220
Met Ser Ser Val Ser Leu Ile Glu Phe Pro Leu Tyr Met Ile Cys Pro
1 5 10 15
Phe Ala Leu Ala Ala Phe Lys Thr Phe Ser Leu Ala Leu Ile Leu Asp
20 25 30
Ile Leu Leu Thr Ile Phe Leu Asp Asp Ile His Phe Val *
35 40 45

<210> 1221
<211> 56
<212> PRT
<213> Homo sapiens

<400> 1221
Met Leu Ile Leu Leu Leu Leu Glu Phe Gly Ile Thr Ile Ile Lys Val
1 5 10 15
Thr Cys Arg Leu Arg Ile Val Leu Cys Tyr Arg Lys Tyr Lys Thr Lys
20 25 30
Arg Asn Lys Lys Leu Lys Leu Gly Asn Asn Ser Lys Phe Gln Arg Met
35 40 45
Cys Leu Arg Thr Ser Phe His *
50 55

<210> 1222
<211> 253
<212> PRT
<213> Homo sapiens

<400> 1222
Met Gly Cys Ala Ile Ile Ala Gly Phe Leu His Tyr Leu Phe Leu Ala
1 5 10 15
Cys Phe Phe Trp Met Leu Val Glu Ala Val Ile Leu Phe Leu Met Val
20 25 30
Arg Asn Leu Lys Val Val Asn Tyr Phe Ser Ser Arg Asn Ile Lys Met
35 40 45
Leu His Ile Cys Ala Phe Gly Tyr Gly Leu Pro Met Leu Val Val Val
50 55 60
Ile Ser Ala Ser Val Gln Pro Gln Gly Tyr Gly Met His Asn Arg Cys
65 70 75 80
Trp Leu Asn Thr Glu Thr Gly Phe Ile Trp Ser Phe Leu Gly Pro Val
85 90 95
Cys Thr Val Ile Val Ile Asn Ser Leu Leu Leu Thr Trp Thr Leu Trp
100 105 110
Ile Leu Arg Gln Arg Leu Ser Ser Val Asn Ala Glu Val Ser Thr Leu

```
        115                 120                 125
Lys Asp Thr Arg Leu Leu Thr Phe Lys Ala Phe Ala Gln Leu Phe Ile
    130                 135                 140
Leu Gly Cys Ser Trp Val Leu Gly Ile Phe Gln Ile Gly Pro Val Ala
145                 150                 155                 160
Gly Val Met Ala Tyr Leu Phe His His His Gln Gln Pro Ala Gly Gly
                165                 170                 175
Leu His Leu Pro His Pro Leu Ser Ala Gln Arg Pro Gly Thr Arg Arg
            180                 185                 190
Ile Gln Glu Val Asp His Trp Glu Asp Glu Ala Gln Leu Pro Val Pro
        195                 200                 205
Asp Leu Lys Asp Leu Ala Val Leu His Ala Ile Arg Phe Gln Asp Gly
    210                 215                 220
Leu Lys Ser Phe Leu Ala Phe Lys Tyr Ala Met Glu Pro Thr Val Gly
225                 230                 235                 240
Gly Thr Ser Ser Phe Pro Cys Arg Glu Pro Tyr Pro  *
                245                 250     252
```

<210> 1223
<211> 858
<212> PRT
<213> Homo sapiens

<400> 1223

```
Met Lys Met Leu Thr Arg Leu Gln Val Leu Thr Leu Ala Leu Phe Ser
 1               5                  10                  15
Lys Gly Phe Leu Leu Ser Leu Gly Asp His Asn Phe Leu Arg Arg Glu
            20                  25                  30
Ile Lys Ile Glu Gly Asp Leu Val Leu Gly Gly Leu Phe Pro Ile Asn
        35                  40                  45
Glu Lys Gly Thr Gly Thr Glu Glu Cys Gly Arg Ile Asn Glu Asp Arg
    50                  55                  60
Gly Ile Gln Arg Leu Glu Ala Met Leu Phe Ala Ile Asp Glu Ile Asn
65                  70                  75                  80
Lys Asp Asp Tyr Leu Leu Pro Gly Val Lys Leu Gly Val His Ile Leu
                85                  90                  95
Asp Thr Cys Ser Arg Asp Thr Tyr Ala Leu Glu Gln Ser Leu Glu Phe
            100                 105                 110
Val Arg Ala Ser Leu Thr Lys Val Asp Glu Ala Glu Tyr Met Cys Pro
        115                 120                 125
Asp Gly Ser Tyr Ala Ile Gln Glu Asn Ile Pro Leu Leu Ile Ala Gly
    130                 135                 140
Val Ile Gly Gly Ser Tyr Ser Arg Val Ser Ile Gln Gly Ala Asn Leu
145                 150                 155                 160
Leu Arg Leu Phe Gln Ile Pro Gln Ile Arg Tyr Ala Ser Thr Ser Ala
                165                 170                 175
Lys Leu Ser Asp Lys Ser Arg Tyr Asp Tyr Phe Ala Arg Thr Val Pro
            180                 185                 190
Pro Asp Phe Tyr Gln Ala Lys Ala Met Ala Glu Ile Leu Arg Phe Phe
        195                 200                 205
Asn Trp Thr Tyr Val Ser Thr Val Ala Ser Glu Gly Asp Tyr Gly Glu
    210                 215                 220
Thr Gly Ile Glu Ala Phe Glu Gln Glu Ala Arg Leu Arg Asn Ile Cys
225                 230                 235                 240
Ile Ala Thr Ala Glu Lys Val Gly Arg Ser Asn Ile Arg Lys Ser Tyr
                245                 250                 255
```

Asp Ser Val Ile Arg Glu Leu Leu Gln Lys Pro Asn Ala Arg Val Val
260 265 270
Val Leu Phe Met Arg Ser Asp Asp Ser Arg Glu Leu Ile Ala Ala Ala
275 280 285
Ser Arg Ala Asn Ala Ser Phe Thr Trp Val Ala Ser Asp Gly Trp Gly
290 295 300
Ala Gln Glu Ser Ile Ile Lys Gly Ser Glu His Val Ala Tyr Gly Ala
305 310 315 320
Ile Thr Leu Glu Leu Ala Ser Gln Pro Val Arg Gln Phe Asp Arg Tyr
325 330 335
Phe Gln Ser Leu Asn Pro Tyr Asn Asn His Arg Asn Pro Trp Phe Arg
340 345 350
Asp Phe Trp Glu Gln Lys Phe Gln Cys Ser Leu Gln Asn Lys Arg Asn
355 360 365
His Arg Arg Val Cys Asp Lys His Leu Ala Ile Asp Ser Ser Asn Tyr
370 375 380
Glu Gln Glu Ser Lys Ile Met Phe Val Val Asn Ala Val Tyr Ala Met
385 390 395 400
Ala His Ala Leu His Lys Met Gln Arg Thr Leu Cys Pro Asn Thr Thr
405 410 415
Lys Leu Cys Asp Ala Met Lys Ile Leu Asp Gly Lys Lys Leu Tyr Lys
420 425 430
Asp Tyr Leu Leu Lys Ile Asn Phe Thr Ala Pro Phe Asn Pro Asn Lys
435 440 445
Asp Ala Asp Ser Ile Val Lys Phe Asp Thr Phe Gly Asp Gly Met Gly
450 455 460
Arg Tyr Asn Val Phe Asn Phe Gln Asn Val Gly Gly Lys Tyr Ser Tyr
465 470 475 480
Leu Lys Val Gly His Trp Ala Glu Thr Leu Ser Leu Asp Val Asn Ser
485 490 495
Ile His Trp Ser Arg Asn Ser Val Pro Thr Ser Gln Cys Ser Asp Pro
500 505 510
Cys Ala Pro Asn Glu Met Lys Asn Met Gln Pro Gly Asp Val Cys Cys
515 520 525
Trp Ile Cys Ile Pro Cys Glu Pro Tyr Glu Tyr Leu Ala Asp Glu Phe
530 535 540
Thr Cys Met Asp Cys Gly Ser Gly Gln Trp Pro Thr Ala Asp Leu Thr
545 550 555 560
Gly Cys Tyr Asp Leu Pro Glu Asp Tyr Ile Arg Trp Glu Asp Ala Trp
565 570 575
Ala Ile Gly Pro Val Thr Ile Ala Cys Leu Gly Phe Met Cys Thr Cys
580 585 590
Met Val Val Thr Val Phe Ile Lys His Asn Asn Thr Pro Leu Val Lys
595 600 605
Ala Ser Gly Arg Glu Leu Cys Tyr Ile Leu Leu Phe Gly Val Gly Leu
610 615 620
Ser Tyr Cys Met Thr Phe Phe Phe Ile Ala Lys Pro Ser Pro Val Ile
625 630 635 640
Cys Ala Leu Arg Arg Leu Gly Leu Gly Ser Ser Phe Ala Ile Cys Tyr
645 650 655
Ser Ala Leu Leu Thr Lys Thr Asn Cys Ile Ala Arg Ile Phe Asp Gly
660 665 670
Val Lys Asn Gly Ala Gln Arg Pro Lys Phe Ile Ser Pro Ser Ser Gln
675 680 685
Val Phe Ile Cys Leu Gly Leu Ile Leu Val Gln Ile Val Met Val Ser
690 695 700
Val Trp Leu Ile Leu Glu Ala Pro Gly Thr Arg Arg Tyr Thr Leu Ala
705 710 715 720
Glu Lys Arg Glu Thr Val Ile Leu Lys Cys Asn Val Lys Asp Ser Ser

```
                725                 730                 735
Met Leu Ile Ser Leu Thr Tyr Asp Val Ile Leu Val Ile Leu Cys Thr
            740                 745                 750
Val Tyr Ala Phe Lys Thr Arg Lys Cys Pro Glu Asn Phe Asn Glu Ala
        755                 760                 765
Lys Phe Ile Gly Phe Thr Met Tyr Thr Thr Cys Ile Ile Trp Leu Ala
    770                 775                 780
Phe Leu Pro Ile Phe Tyr Val Thr Ser Ser Asp Tyr Arg Val Gln Thr
785                 790                 795                 800
Thr Thr Met Cys Ile Ser Val Ser Leu Ser Gly Phe Val Val Leu Gly
                805                 810                 815
Cys Leu Phe Ala Pro Lys Val His Ile Ile Leu Phe Gln Pro Gln Lys
            820                 825                 830
Asn Val Val Thr His Arg Leu His Leu Asn Arg Phe Ser Val Ser Gly
        835                 840                 845
Thr Gly Thr His Ile Leu Ser Val Leu  *
    850                 855     857
```

<210> 1224
<211> 69
<212> PRT
<213> Homo sapiens

<400> 1224

```
Met Ser His Met Val Pro Leu Ala Leu Leu Leu Pro Leu Phe Pro Thr
 1               5                  10                  15
Ser Arg Arg Ala Ala Leu Pro Phe Leu Pro Leu Phe Phe Gly Leu Met
            20                  25                  30
Phe Pro Ala Thr Thr Asp Leu Pro Pro Pro His Pro Ser Ala Asp Leu
        35                  40                  45
Ala Val His Cys Arg His Gly Gly Leu Ile Ser Asp Arg Lys Leu Arg
    50                  55                  60
Leu Ser Glu Arg  *
 65         68
```

<210> 1225
<211> 55
<212> PRT
<213> Homo sapiens

<400> 1225

```
Met Cys Tyr His Thr Trp Leu Ile Phe Ile Phe Leu Val Glu Met Gly
 1               5                  10                  15
Phe Tyr His Val Gly Gln Ala Gly Phe Lys Leu Leu Ala Ser Ser Gly
            20                  25                  30
Pro Pro Ala Ser Ala Ser Gln Ser Ala Gly Ile Thr Gly Val Ser His
        35                  40                  45
His Ala Arg Pro Thr Phe  *
    50              54
```

<210> 1226

<211> 51
<212> PRT
<213> Homo sapiens

<400> 1226
Met Ile Leu Ser Leu Leu Lys Phe Phe Pro Leu Leu Ser Ser Asp Thr
1 5 10 15
Pro Asn Ser Ser Val Pro Leu Leu Thr Thr Pro Arg Asp Pro Pro Tyr
20 25 30
His Leu Ser Pro Cys Ser Ser Ser Tyr Phe Val Lys Glu Gly Phe Ser
35 40 45
Val Val *
50

<210> 1227
<211> 47
<212> PRT
<213> Homo sapiens

<400> 1227
Met Ile Leu Phe Cys Val Met Val Phe Ile Leu Phe Ile Thr Phe His
1 5 10 15
Leu Gln Leu Pro Thr Val Gly Asp Val Thr Tyr Cys Phe Cys Ser Asn
20 25 30
Lys Leu Arg Lys Thr Arg Glu Leu Lys Lys Ile Ser Ser Asn *
35 40 45 46

<210> 1228
<211> 60
<212> PRT
<213> Homo sapiens

<400> 1228
Met Phe Ser Thr Ala Phe Trp Pro Pro Phe Leu Asn Pro Ser Leu Met
1 5 10 15
Phe Phe Thr Leu Leu Cys Ser Asp Phe Met Pro Cys Glu Ala Val Cys
20 25 30
Ser Ser Ile Ile Tyr Ser Phe Ile Pro Val Thr Lys Thr Gln Gly Ala
35 40 45
Ala Pro His Thr Arg Gly Pro Gln Pro His Thr *
50 55 59

<210> 1229
<211> 52
<212> PRT
<213> Homo sapiens

<400> 1229
Met Cys Glu Ser Thr Glu Leu Asn Met Thr Phe His Leu Phe Ile Val

```
 1               5                   10                  15
Ala Leu Ala Gly Ala Gly Ala Ala Val Ile Ala Met Val His Tyr Leu
            20                  25                  30
Met Val Leu Ser Ala Asn Trp Ala Tyr Val Lys Asp Ala Cys Arg Met
        35                  40                  45
Ala Glu Val  *
    50  51
```

<210> 1230
<211> 362
<212> PRT
<213> Homo sapiens

<400> 1230

```
Met Pro Val Ile Trp Ser Ala Leu Ser Ala Val Leu Leu Leu Ala Ser
 1               5                   10                  15
Ser Tyr Phe Val Gly Ala Leu Ile Val His Ala Asp Cys Phe Leu Met
            20                  25                  30
Arg Asn His Thr Ile Thr Glu Gln Pro Met Cys Phe Gln Arg Thr Thr
        35                  40                  45
Pro Leu Ile Leu Gln Glu Val Ala Ser Phe Leu Lys Arg Asn Lys His
    50                  55                  60
Gly Pro Phe Leu Leu Phe Val Ser Phe Leu His Val His Ile Pro Leu
65                  70                  75                  80
Ile Thr Met Glu Asn Phe Leu Gly Lys Ser Leu His Gly Leu Tyr Gly
                85                  90                  95
Asp Asn Val Lys Glu Met Asp Trp Met Val Gly Arg Ile Leu Asp Thr
            100                 105                 110
Leu Asp Val Glu Gly Leu Ser Asn Ser Thr Leu Ile Tyr Phe Thr Ser
        115                 120                 125
Asp His Gly Gly Ser Leu Glu Asn Gln Leu Gly Asn Thr Gln Tyr Gly
    130                 135                 140
Gly Trp Asn Gly Ile Tyr Lys Gly Gly Lys Gly Met Gly Gly Trp Glu
145                 150                 155                 160
Gly Gly Ile Arg Val Pro Gly Ile Phe Arg Trp Pro Gly Val Leu Pro
                165                 170                 175
Ala Gly Arg Val Ile Gly Glu Pro Thr Ser Leu Met Asp Val Phe Pro
            180                 185                 190
Thr Val Val Arg Leu Ala Gly Ser Glu Val Pro Gln Asp Arg Val Ile
        195                 200                 205
Asp Gly Gln Asp Leu Leu Pro Leu Leu Leu Gly Thr Ala Gln His Ser
    210                 215                 220
Asp His Glu Phe Leu Met His Tyr Cys Glu Arg Phe Leu His Ala Ala
225                 230                 235                 240
Arg Trp His Gln Arg Asp Arg Gly Thr Met Trp Lys Val His Phe Val
                245                 250                 255
Thr Pro Val Phe Gln Pro Arg Gly Ser Arg Cys Leu Leu Trp Lys Glu
            260                 265                 270
Lys Val Cys Pro Cys Phe Gly Glu Lys Ser Ser Pro Pro Arg Ser His
        275                 280                 285
Pro Cys Phe Phe Asp Leu Ser Arg Ala Pro Ser Glu Thr His Ile Leu
    290                 295                 300
Thr Pro Ala Ser Glu Pro Val Phe Tyr Gln Val Met Glu Arg Ser Pro
305                 310                 315                 320
Ala Gly Gly Val Gly Thr Pro Ala Asp Thr Gln Pro Ser Ser Ser Ala
                325                 330                 335
```

```
Ala Gly Gln Ala Gly Gln Tyr Leu Glu Thr Gly Gly Ala Ala Leu Leu
            340                 345                 350
Trp Ala Val Pro Pro Leu Val Gly Pro  *
        355                 360 361
```

<210> 1231
<211> 53
<212> PRT
<213> Homo sapiens

<400> 1231

```
Met Leu Arg Leu Gly Val Ala Phe His Met Glu Leu Leu Cys Arg Gly
 1               5                  10                  15
Arg Leu Leu Leu Leu Ile Pro Thr Ala Glu Thr Arg Cys Asp His Arg
            20                  25                  30
Arg Leu Gln Asn Leu Lys Leu Gly Leu Ser Asn Thr Leu Asp Lys His
        35                  40                  45
Gln Glu Pro His  *
    50      52
```

<210> 1232
<211> 56
<212> PRT
<213> Homo sapiens

<400> 1232

```
Met Leu Asn Phe Ile Ser Pro Phe Gly Ser Thr Ile Leu Leu Leu Ile
 1               5                  10                  15
Pro Ser Ala Leu Pro Pro Ser Pro Pro Ser Arg Cys Ser Leu Leu Ser
            20                  25                  30
Pro Pro Pro Thr Thr Pro Leu Pro Leu Pro Leu Pro Ser Pro Phe Ser
        35                  40                  45
Ser Pro Leu Leu Ser Phe Phe  *
    50                  55
```

<210> 1233
<211> 56
<212> PRT
<213> Homo sapiens

<400> 1233

```
Met Gln Leu His Val Ser Leu Pro Trp Leu Leu Arg Phe Pro Gly Leu
 1               5                  10                  15
Asp Cys Thr Leu His Pro Asp Gln Pro Ser Ile Gln Leu Leu Gln Gly
            20                  25                  30
Thr Ile Asp Leu Leu Asp Ser Val Ile Leu Ser Cys Ser Leu Cys Leu
        35                  40                  45
Phe Gly Val Leu Gln Met His Ile
    50                  55  56
```

<210> 1234
<211> 125
<212> PRT
<213> Homo sapiens

<400> 1234
Met Leu Ser Gln Leu Pro Arg Cys Gln Ser Ser Val Pro Ala Leu Ala
1 5 10 15
His Pro Thr Arg Leu His Tyr Leu Leu Arg Leu Leu Thr Phe Leu Leu
20 25 30
Gly Pro Gly Ala Gly Gly Ala Glu Ala Gln Gly Met Leu Gly Arg Ala
35 40 45
Leu Leu Leu Ser Ser Leu Pro Asp Asn Cys Ser Phe Trp Asp Ala Phe
50 55 60
Arg Pro Glu Gly Arg Arg Ser Val Leu Arg Thr Ile Gly Glu Tyr Leu
65 70 75 80
Glu Gln Asp Glu Glu Gln Pro Thr Pro Ser Gly Phe Glu Pro Thr Val
85 90 95
Asn Pro Ser Ser Gly Ile Ser Lys Met Glu Leu Leu Ala Cys Phe Ser
100 105 110
Val Ser Ala Leu Pro Glu Gly Lys Leu Leu Glu Gln *
115 120 124

<210> 1235
<211> 72
<212> PRT
<213> Homo sapiens

<400> 1235
Met Phe Cys Phe Leu His Val Phe Leu Val Ser Leu Pro Phe Leu Thr
1 5 10 15
Ser Tyr Ser Cys Leu Gln Ile Ile Ser Tyr Ser Ser Phe Lys Ala Trp
20 25 30
Phe Lys Tyr Pro Phe Leu Cys Lys Ile Phe Pro Thr Leu Pro Asn Asn
35 40 45
Asp Ser Leu Gln Gln Thr Pro Leu Val His Gly Val Cys Leu Gln Gln
50 55 60
Gly Val His His Arg Leu Ile *
65 70 71

<210> 1236
<211> 48
<212> PRT
<213> Homo sapiens

<400> 1236
Met Ala Pro Gly Gly Ala Lys Gly Gln Gly Ala Ser Ala Leu Ala Leu
1 5 10 15
Leu Phe Ile Leu Ala Ser Pro Ala Thr Gly Gly Gly Pro Arg Leu Trp
20 25 30

```
Arg Ala Gly Gly Leu Gly Phe Thr His Cys Gln Ala Asn Ser Thr Thr
        35                  40                  45          48
```

<210> 1237
<211> 208
<212> PRT
<213> Homo sapiens

<400> 1237

```
Met Ala Phe Leu Arg Lys Val Tyr Ser Ile Leu Ser Leu Gln Val Leu
  1               5                  10                  15
Leu Thr Thr Val Thr Ser Thr Val Phe Leu Tyr Phe Glu Ser Val Arg
             20                  25                  30
Thr Phe Val His Glu Ser Pro Ala Leu Ile Leu Leu Phe Ala Leu Gly
        35                  40                  45
Ser Leu Gly Leu Ile Phe Ala Leu Ile Leu Asn Arg His Lys Tyr Pro
    50                  55                  60
Leu Asn Leu Tyr Leu Leu Phe Gly Phe Thr Leu Leu Glu Ala Leu Thr
 65                  70                  75                      80
Val Ala Val Val Val Thr Phe Tyr Asp Val Tyr Ile Ile Leu Gln Ala
                 85                  90                  95
Phe Ile Leu Thr Thr Thr Val Phe Phe Gly Leu Thr Val Tyr Thr Leu
            100                 105                 110
Gln Ser Lys Lys Asp Phe Ser Lys Phe Gly Ala Gly Leu Phe Ala Leu
        115                 120                 125
Leu Trp Ile Leu Cys Leu Ser Gly Phe Leu Lys Phe Phe Phe Tyr Ser
    130                 135                 140
Glu Ile Met Glu Leu Val Leu Ala Ala Ala Gly Ala Leu Leu Phe Cys
145                 150                 155                     160
Gly Phe Ile Ile Tyr Asp Thr His Ser Leu Met His Lys Leu Ser Pro
                165                 170                 175
Glu Glu Tyr Val Leu Ala Ala Ile Ser Leu Tyr Leu Asp Ile Ile Asn
            180                 185                 190
Leu Phe Leu His Leu Leu Arg Phe Leu Glu Ala Val Asn Lys Lys  *
        195                 200                 205     207
```

<210> 1238
<211> 173
<212> PRT
<213> Homo sapiens

<400> 1238

```
Met Lys Val Val Pro Ser Leu Leu Leu Ser Val Leu Leu Ala Gln Val
  1               5                  10                  15
Trp Leu Val Pro Gly Leu Ala Pro Ser Pro Gln Ser Pro Glu Thr Pro
             20                  25                  30
Ala Pro Gln Asn Gln Thr Ser Arg Val Val Gln Ala Pro Lys Glu Glu
        35                  40                  45
Glu Glu Asp Glu Gln Glu Ala Ser Glu Glu Lys Ala Ser Glu Glu Glu
    50                  55                  60
Lys Ala Trp Leu Met Ala Ser Arg Gln Gln Leu Ala Lys Glu Thr Ser
```

```
65                  70                  75                  80
Asn Phe Gly Phe Ser Leu Leu Arg Lys Ile Ser Met Arg His Asp Gly
                85                  90                  95
Asn Met Val Phe Ser Pro Phe Gly Met Ser Leu Ala Met Thr Gly Leu
            100                 105                 110
Met Leu Gly Ala Thr Gly Pro Thr Glu Thr Gln Ile Lys Arg Gly Leu
        115                 120                 125
His Leu Gln Ala Leu Lys Pro Thr Lys Pro Gly Leu Leu Pro Ser Leu
    130                 135                 140
Phe Lys Gly Leu Arg Glu Thr Leu Ser Arg Asn Leu Glu Leu Gly Leu
145                 150                 155                 160
Thr Ala Gly Glu Phe Cys Leu His Pro Gln Gly Phe  *
                165                 170     172
```

<210> 1239
<211> 357
<212> PRT
<213> Homo sapiens

<400> 1239

```
Met Ala Phe Leu Gly Leu Phe Ser Leu Leu Val Leu Gln Ser Met Ala
1               5                   10                  15
Thr Gly Ala Thr Phe Pro Glu Glu Ala Ile Ala Asp Leu Ser Val Asn
            20                  25                  30
Met Tyr Asn Arg Leu Arg Ala Thr Gly Glu Asp Glu Asn Ile Leu Phe
        35                  40                  45
Ser Pro Leu Ser Ile Ala Leu Ala Met Gly Met Met Glu Leu Gly Ala
    50                  55                  60
Gln Gly Ser Thr Gln Lys Glu Ile Arg His Ser Met Gly Tyr Asp Ser
65                  70                  75                  80
Leu Lys Asn Gly Glu Glu Phe Ser Phe Leu Lys Glu Phe Ser Asn Met
                85                  90                  95
Val Thr Ala Lys Glu Ser Gln Tyr Val Met Lys Ile Ala Asn Ser Leu
            100                 105                 110
Phe Val Gln Asn Gly Phe His Val Asn Glu Glu Phe Leu Gln Met Met
        115                 120                 125
Lys Lys Tyr Phe Asn Ala Ala Val Asn His Val Asp Phe Ser Gln Asn
    130                 135                 140
Val Ala Val Ala Asn Tyr Ile Asn Lys Trp Val Glu Asn Asn Thr Asn
145                 150                 155                 160
Asn Leu Val Lys Asp Leu Val Ser Pro Arg Asp Phe Asp Ala Ala Thr
                165                 170                 175
Tyr Leu Ala Leu Ile Asn Ala Val Tyr Phe Lys Gly Asn Trp Lys Ser
            180                 185                 190
Gln Phe Arg Pro Glu Asn Thr Arg Thr Phe Ser Phe Thr Lys Asp Asp
        195                 200                 205
Glu Ser Glu Val Gln Ile Pro Met Met Tyr Gln Gln Gly Glu Phe Tyr
    210                 215                 220
Tyr Gly Glu Phe Ser Asp Gly Ser Asn Glu Ala Gly Gly Ile Tyr Gln
225                 230                 235                 240
Val Leu Glu Ile Pro Tyr Glu Gly Asp Glu Ile Ser Met Met Leu Val
                245                 250                 255
Leu Ser Arg Gln Glu Val Pro Leu Ala Thr Leu Glu Pro Leu Val Lys
            260                 265                 270
Ala Gln Leu Val Glu Glu Trp Ala Asn Ser Val Lys Lys Gln Lys Val
        275                 280                 285
```

```
Glu Val Tyr Leu Pro Arg Phe Thr Val Glu Gln Glu Ile Asp Leu Lys
    290                 295                 300
Asp Val Leu Lys Ala Leu Gly Ile Thr Glu Ile Phe Ile Lys Asp Ala
305                 310                 315                 320
Asn Leu Thr Gly Leu Ser Asp Asn Lys Glu Ile Phe Leu Ser Lys Ala
                325                 330                 335
Ile His Lys Ser Phe Leu Glu Val Asn Glu Glu Ala Gln Lys Leu Leu
            340                 345                 350
Leu Ser Gln Glu  *
        355 356
```

<210> 1240
<211> 707
<212> PRT
<213> Homo sapiens

<400> 1240

```
Met Leu Ser Leu Arg Arg Cys Thr Ser Met Arg Leu Cys Leu Ser Ser
  1               5                  10                  15
Ser Leu Ala Ser Pro Cys Ser Thr Met Leu Ser Thr Val Val Leu Tyr
            20                  25                  30
Lys Val Cys Asn Ser Phe Val Glu Met Gly Ser Ala Asn Val Gln Ala
        35                  40                  45
Thr Asp Tyr Leu Lys Gly Val Ala Ser Leu Phe Val Val Ser Leu Gly
    50                  55                  60
Gly Ala Ala Val Gly Leu Val Phe Ala Phe Leu Leu Ala Leu Thr Thr
 65                  70                  75                  80
Arg Phe Thr Lys Arg Val Arg Ile Ile Glu Pro Leu Leu Val Phe Leu
                85                  90                  95
Leu Ala Tyr Ala Ala Tyr Leu Thr Ala Glu Met Ala Ser Leu Ser Ala
            100                 105                 110
Ile Leu Ala Val Thr Met Cys Gly Leu Gly Cys Lys Lys Tyr Val Glu
        115                 120                 125
Ala Asn Ile Ser His Lys Ser Arg Thr Thr Val Lys Tyr Thr Met Lys
    130                 135                 140
Thr Leu Ala Ser Cys Ala Glu Thr Val Ile Phe Met Leu Leu Gly Ile
145                 150                 155                 160
Ser Thr Val Asp Ser Ser Lys Trp Ala Trp Asp Ser Gly Leu Val Leu
                165                 170                 175
Gly Thr Leu Ile Phe Ile Leu Phe Phe Arg Ala Leu Gly Val Val Leu
            180                 185                 190
Gln Thr Trp Val Leu Asn Gln Phe Arg Leu Val Pro Leu Asp Lys Ile
        195                 200                 205
Asp Gln Val Val Met Ser Tyr Gly Gly Leu Arg Gly Ala Val Ala Phe
    210                 215                 220
Ala Leu Val Ile Leu Leu Asp Arg Thr Lys Val Pro Ala Lys Asp Tyr
225                 230                 235                 240
Phe Val Ala Thr Thr Ile Val Val Val Phe Phe Thr Val Ile Val Gln
                245                 250                 255
Gly Leu Thr Ile Lys Pro Leu Val Lys Trp Leu Lys Val Lys Arg Ser
            260                 265                 270
Glu His His Lys Pro Thr Leu Asn Gln Glu Leu His Glu His Thr Phe
        275                 280                 285
Asp His Ile Leu Ala Ala Val Glu Asp Val Val Gly His His Gly Tyr
    290                 295                 300
His Tyr Trp Arg Asp Arg Trp Glu Gln Phe Asp Lys Lys Tyr Leu Ser
```

305 310 315 320
Gln Leu Leu Met Arg Arg Ser Ala Tyr Arg Ile Arg Asp Gln Ile Trp
325 330 335
Asp Val Tyr Tyr Arg Leu Asn Ile Arg Asp Ala Ile Ser Phe Val Asp
340 345 350
Gln Gly Gly His Val Leu Ser Ser Thr Gly Leu Thr Leu Pro Ser Met
355 360 365
Pro Ser Arg Asn Ser Val Ala Glu Thr Ser Val Thr Asn Leu Leu Arg
370 375 380
Glu Ser Gly Ser Gly Ala Cys Leu Asp Leu Gln Val Ile Asp Thr Val
385 390 395 400
Arg Ser Gly Arg Asp Arg Glu Asp Ala Val Met His His Leu Leu Cys
405 410 415
Gly Gly Leu Tyr Lys Pro Arg Arg Arg Tyr Lys Ala Ser Cys Ser Arg
420 425 430
His Phe Ile Ser Glu Asp Ala Gln Glu Arg Gln Asp Lys Glu Val Phe
435 440 445
Gln Gln Asn Met Lys Arg Arg Leu Glu Ser Phe Lys Ser Thr Lys His
450 455 460
Asn Ile Cys Phe Thr Lys Ser Lys Pro Arg Pro Arg Lys Thr Gly Arg
465 470 475 480
Arg Lys Lys Asp Gly Val Ala Asn Ala Glu Ala Thr Asn Gly Lys His
485 490 495
Arg Gly Leu Gly Phe Gln Asp Thr Ala Ala Val Ile Leu Thr Val Glu
500 505 510
Ser Glu Glu Glu Glu Glu Glu Ser Asp Ser Ser Glu Thr Glu Lys Glu
515 520 525
Asp Asp Glu Gly Ile Ile Phe Val Ala Arg Ala Thr Ser Glu Val Leu
530 535 540
Gln Glu Gly Lys Val Ser Gly Ser Leu Glu Val Cys Pro Ser Pro Arg
545 550 555 560
Ile Ile Pro Pro Ser Pro Thr Cys Ala Glu Lys Glu Leu Pro Trp Lys
565 570 575
Ser Gly Gln Gly Asp Leu Ala Val Tyr Val Ser Ser Glu Thr Thr Lys
580 585 590
Ile Val Pro Val Asp Met Gln Thr Gly Trp Asn Gln Ser Ile Ser Ser
595 600 605
Leu Glu Ser Leu Ala Ser Pro Pro Cys Asn Gln Ala Pro Ile Leu Thr
610 615 620
Cys Leu Pro Pro His Pro Arg Gly Thr Glu Glu Pro Gln Val Pro Leu
625 630 635 640
His Leu Pro Ser Asp Pro Arg Ser Ser Phe Ala Phe Pro Pro Ser Leu
645 650 655
Ala Lys Ala Gly Arg Ser Arg Ser Glu Ser Ser Ala Asp Leu Pro Gln
660 665 670
Gln Gln Glu Leu Gln Pro Leu Met Gly His Lys Asp His Thr His Leu
675 680 685
Ser Pro Gly Thr Ala Thr Ser His Trp Cys Ile Gln Phe Asn Arg Gly
690 695 700
Ser Arg Leu
705 707

<210> 1241
<211> 98
<212> PRT
<213> Homo sapiens

<400> 1241
Met Ala Phe Arg Thr Phe Ser Trp Ile Phe Ser Gly Leu Leu Ser Pro
1 5 10 15
Thr Leu Ala Ser Pro Ser Val Ser Met Met Thr Met Glu Val Leu Leu
20 25 30
Ser Gly Ile Leu Cys Ser Ser Arg Ala Leu Phe Ser Ile Leu Met Pro
35 40 45
Leu Ser Ser Pro Ser Leu Met Leu Val Ile Pro Leu Ser Ser Met Leu
50 55 60
Phe Thr Asn Val Leu Ala Ser Trp Arg Phe Ser Gly Val Ala Trp Thr
65 70 75 80
Lys Cys Ser Phe His Val Asp Thr Ser Pro Leu Asn Arg Met Lys Phe
85 90 95
Arg *
97

<210> 1242
<211> 422
<212> PRT
<213> Homo sapiens

<400> 1242
Met Val Leu Trp Glu Ser Pro Arg Gln Cys Ser Ser Trp Thr Leu Cys
1 5 10 15
Glu Gly Phe Cys Trp Leu Leu Leu Leu Pro Val Met Leu Leu Ile Val
20 25 30
Ala Arg Pro Val Lys Leu Ala Ala Phe Pro Thr Ser Leu Ser Asp Cys
35 40 45
Gln Thr Pro Thr Gly Trp Asn Cys Ser Gly Tyr Asp Asp Arg Glu Asn
50 55 60
Asp Leu Phe Leu Cys Asp Thr Asn Thr Cys Lys Phe Asp Gly Glu Cys
65 70 75 80
Leu Arg Ile Gly Asp Thr Val Thr Cys Val Cys Gln Phe Lys Cys Asn
85 90 95
Asn Asp Tyr Val Pro Val Cys Gly Ser Asn Gly Glu Ser Tyr Gln Asn
100 105 110
Glu Cys Tyr Leu Arg Gln Ala Ala Cys Lys Gln Gln Ser Glu Ile Leu
115 120 125
Val Val Ser Glu Gly Ser Cys Ala Thr Asp Ala Gly Ser Gly Ser Gly
130 135 140
Asp Gly Val His Glu Gly Ser Gly Glu Thr Ser Gln Lys Glu Thr Ser
145 150 155 160
Thr Cys Asp Ile Cys Gln Phe Gly Ala Glu Cys Asp Glu Asn Ala Glu
165 170 175
Asp Val Trp Cys Val Cys Asn Ile Asp Cys Ser Gln Thr Asn Phe Asn
180 185 190
Pro Leu Cys Ala Ser Asp Gly Lys Ser Tyr Asp Asn Ala Cys Gln Ile
195 200 205
Lys Glu Ala Ser Cys Gln Lys Gln Glu Lys Ile Glu Val Leu Ser Leu
210 215 220
Gly Arg Cys Gln Asp Asn Thr Thr Thr Thr Thr Lys Ser Glu Asp Gly
225 230 235 240
His Tyr Ala Arg Thr Asp Tyr Ala Glu Asn Ala Asn Lys Leu Glu Glu
245 250 255
Ser Ala Arg Glu His His Ile Pro Cys Pro Glu His Tyr Asn Gly Phe

```
            260                 265                 270
Cys Met His Gly Lys Cys Glu His Ser Ile Asn Met Gln Glu Pro Ser
        275                 280                 285
Cys Arg Cys Asp Ala Gly Tyr Thr Gly Gln His Cys Glu Lys Lys Asp
    290                 295                 300
Tyr Ser Val Leu Tyr Val Val Pro Gly Pro Val Arg Phe Pro Val Cys
305                 310                 315                 320
Leu Asn Arg Ser Cys Asp Trp Asn Asn Ser Asp Cys Cys His Leu Cys
                325                 330                 335
Gly Gly Pro Leu His His Lys Glu Met Pro Pro Glu Ala Asn Arg Ile
            340                 345                 350
Pro Pro Asp Arg Ser Lys Ile Pro Gly His Tyr Ser Ser Arg Gln Tyr
        355                 360                 365
Asn Lys Ser Arg Pro Thr Arg Leu Ile Leu Lys Gly Ala Cys Phe His
    370                 375                 380
Ser Gly Trp Thr Thr Glu Ser Leu Asp Tyr Thr Ile Gln Tyr Tyr Arg
385                 390                 395                 400
Gln Lys Asn Lys Thr Arg Asp Leu Thr His Val Cys Leu Ala Phe Val
                405                 410                 415
Gly Asn Leu His Gln  *
            420 421
```

<210> 1243
<211> 46
<212> PRT
<213> Homo sapiens

<400> 1243

```
Met Leu Phe Val Phe Ile Cys Ser Tyr Phe His Leu Ser Leu Phe Leu
 1               5                  10                  15
Leu Phe Pro Phe Leu Pro Val Ser Leu Pro Ser Phe Leu Pro Phe Phe
            20                  25                  30
Leu Pro Ser Phe Leu Glu Phe Thr Glu Val Phe Pro Arg  *
        35                  40                  45
```

<210> 1244
<211> 46
<212> PRT
<213> Homo sapiens

<400> 1244

```
Met Val Leu Ser Ala Pro Ser Leu Trp Pro Cys Ser Ser Phe Ser Ile
 1               5                  10                  15
Ser Cys Leu His Val Gly Leu Thr Ala Phe Leu Phe Gln Val Ala Phe
            20                  25                  30
Leu Cys Leu Leu Cys Cys Val Glu Leu Leu Leu Asp Val  *
        35                  40                  45
```

<210> 1245
<211> 244
<212> PRT

<213> Homo sapiens

<400> 1245
Met Ala Gly Val Ile Ala Gly Leu Leu Met Phe Ile Ile Ile Leu Leu
1 5 10 15
Gly Val Met Leu Thr Ile Lys Arg Arg Arg Asn Ala Tyr Ser Tyr Ser
20 25 30
Tyr Tyr Leu Lys Leu Ala Lys Lys Gln Lys Glu Thr Gln Ser Gly Ala
35 40 45
Gln Arg Glu Met Gly Pro Val Ala Ser Ala Asp Lys Pro Thr Thr Lys
50 55 60
Leu Ser Ala Ser Arg Asn Asp Glu Gly Phe Ser Ser Ser Ser Gln Asp
65 70 75 80
Val Asn Gly Phe Asn Gly Ser Arg Gly Glu Leu Ser Gln Pro Thr Leu
85 90 95
Thr Ile Gln Thr His Pro Tyr Arg Thr Cys Asp Pro Val Glu Met Ser
100 105 110
Tyr Pro Arg Asp Gln Phe Gln Pro Ala Ile Arg Val Ala Asp Leu Leu
115 120 125
Gln His Ile Thr Gln Met Lys Arg Gly Gln Gly Tyr Gly Phe Lys Glu
130 135 140
Glu Tyr Glu Ala Leu Pro Glu Gly Gln Thr Ala Ser Trp Asp Thr Ala
145 150 155 160
Lys Glu Asp Glu Asn Arg Asn Lys Asn Arg Tyr Gly Asn Ile Ile Ser
165 170 175
Tyr Asp His Ser Arg Val Arg Leu Leu Val Leu Asp Gly Asp Pro His
180 185 190
Ser Asp Tyr Ile Asn Ala Asn Tyr Ile Asp Gly Tyr His Arg Pro Arg
195 200 205
His Tyr Ile Ala Thr Gln Gly Pro Met Gln Glu Thr Val Lys Asp Phe
210 215 220
Trp Arg Met Ile Trp Gln Glu Asn Ser Ala Ser Ile Val Met Val Thr
225 230 235 240
Asn Pro Gly *
243

<210> 1246
<211> 565
<212> PRT
<213> Homo sapiens

<400> 1246
Met Ala Val Phe Arg Ser Gly Leu Leu Val Leu Thr Thr Pro Leu Ala
1 5 10 15
Ser Leu Ala Pro Arg Leu Ala Ser Ile Leu Thr Ser Ala Ala Arg Leu
20 25 30
Val Asn His Thr Leu Tyr Val His Leu Gln Pro Gly Met Ser Leu Glu
35 40 45
Gly Pro Ala Gln Pro Gln Tyr Ser Pro Val Gln Ala Thr Phe Glu Val
50 55 60
Leu Asp Phe Ile Thr His Leu Tyr Ala Gly Ala Asp Val His Arg His
65 70 75 80
Leu Asp Val Arg Ile Leu Leu Thr Asn Ile Arg Thr Lys Ser Thr Phe
85 90 95
Leu Pro Pro Leu Pro Thr Ser Val Gln Asn Leu Ala His Pro Pro Glu

```
            100                 105                 110
Val Val Leu Thr Asp Phe Gln Thr Leu Asp Gly Ser Gln Tyr Asn Pro
        115                 120                 125
Val Lys Gln Gln Leu Val Arg Tyr Ala Thr Ser Cys Tyr Ser Cys Cys
    130                 135                 140
Pro Arg Leu Ala Ser Val Leu Leu Tyr Ser Asp Tyr Gly Ile Gly Glu
145                 150                 155                 160
Val Pro Val Glu Pro Leu Asp Val Pro Leu Pro Ser Thr Ile Arg Pro
                165                 170                 175
Ala Ser Pro Val Ala Gly Ser Pro Lys Gln Pro Val Arg Gly Tyr Tyr
            180                 185                 190
Arg Gly Ala Val Gly Gly Thr Phe Asp Arg Leu His Asn Ala His Lys
        195                 200                 205
Val Leu Leu Ser Val Ala Cys Ile Leu Ala Gln Glu Gln Leu Val Val
    210                 215                 220
Gly Val Ala Asp Lys Asp Leu Leu Lys Ser Lys Leu Leu Pro Glu Leu
225                 230                 235                 240
Leu Gln Pro Tyr Thr Glu Arg Val Glu His Leu Ser Glu Phe Leu Val
                245                 250                 255
Asp Ile Lys Pro Ser Leu Thr Phe Asp Val Ile Pro Leu Leu Asp Pro
            260                 265                 270
Tyr Gly Pro Ala Gly Ser Asp Pro Ser Leu Glu Phe Leu Val Val Ser
        275                 280                 285
Glu Glu Thr Tyr Arg Gly Gly Met Ala Ile Asn Arg Phe Arg Leu Glu
    290                 295                 300
Asn Asp Leu Glu Glu Leu Ala Leu Tyr Gln Ile Gln Leu Leu Lys Asp
305                 310                 315                 320
Leu Arg His Thr Glu Asn Glu Glu Asp Lys Val Ser Ser Ser Ser Phe
                325                 330                 335
Arg Gln Arg Met Leu Gly Asn Leu Leu Arg Pro Pro Tyr Glu Arg Pro
            340                 345                 350
Glu Leu Pro Thr Cys Leu Tyr Val Ile Gly Leu Thr Gly Ile Ser Gly
        355                 360                 365
Ser Gly Lys Ser Ser Ile Ala Gln Arg Leu Lys Gly Leu Gly Ala Phe
    370                 375                 380
Val Ile Asp Ser Asp His Leu Gly His Arg Ala Tyr Ala Pro Gly Gly
385                 390                 395                 400
Pro Ala Tyr Gln Pro Val Val Glu Ala Phe Gly Thr Asp Ile Leu His
                405                 410                 415
Lys Asp Gly Ile Ile Asn Arg Lys Val Leu Gly Ser Arg Val Phe Gly
            420                 425                 430
Asn Lys Lys Gln Leu Lys Ile Leu Thr Asp Ile Met Trp Pro Ile Ile
        435                 440                 445
Ala Lys Leu Ala Arg Glu Glu Met Asp Arg Ala Val Ala Glu Gly Lys
    450                 455                 460
Arg Val Cys Val Ile Asp Ala Ala Val Leu Leu Glu Ala Gly Trp Gln
465                 470                 475                 480
Asn Leu Val His Glu Val Trp Thr Ala Val Ile Pro Glu Thr Glu Ala
                485                 490                 495
Val Arg Arg Ile Val Glu Arg Asp Gly Leu Ser Glu Ala Ala Ala Gln
            500                 505                 510
Ser Arg Leu Gln Ser Gln Met Ser Gly Gln Gln Leu Val Glu Gln Ser
        515                 520                 525
His Val Val Leu Ser Thr Leu Trp Glu Pro His Ile Thr Gln Arg Gln
    530                 535                 540
Val Glu Lys Ala Trp Ala Leu Leu Gln Lys Arg Ile Pro Lys Thr His
545                 550                 555                 560
Gln Ala Leu Asp  *
            564
```

<210> 1247
<211> 737
<212> PRT
<213> Homo sapiens

<400> 1247
Met Phe Pro Ala Gly Pro Pro Trp Pro Arg Val Arg Val Val Gln Val
1 5 10 15
Leu Trp Ala Leu Leu Ala Val Leu Leu Ala Ser Trp Arg Leu Trp Ala
20 25 30
Ile Lys Asp Phe Gln Glu Cys Thr Trp Gln Val Val Leu Asn Glu Phe
35 40 45
Lys Arg Val Gly Glu Ser Gly Val Ser Asp Ser Phe Phe Glu Gln Glu
50 55 60
Pro Val Asp Thr Val Ser Ser Leu Phe His Met Leu Val Asp Ser Pro
65 70 75 80
Ile Asp Pro Ser Glu Lys Tyr Leu Gly Phe Pro Tyr Tyr Leu Lys Ile
85 90 95
Asn Tyr Ser Cys Glu Glu Lys Pro Ser Glu Asp Leu Val Arg Met Gly
100 105 110
His Leu Thr Gly Leu Lys Pro Leu Val Leu Val Thr Phe Gln Ser Pro
115 120 125
Val Asn Phe Tyr Arg Trp Lys Ile Glu Gln Leu Gln Ile Gln Met Glu
130 135 140
Ala Ala Pro Phe Arg Ser Lys Gly Gly Pro Gly Gly Gly Gly Arg Asp
145 150 155 160
Arg Asn Leu Ala Gly Met Asn Ile Asn Gly Phe Leu Lys Arg Asp Arg
165 170 175
Asp Asn Asn Ile Gln Phe Thr Val Gly Glu Glu Leu Phe Asn Leu Met
180 185 190
Pro Gln Tyr Phe Val Gly Val Ser Ser Arg Pro Leu Trp His Thr Val
195 200 205
Asp Gln Ser Pro Val Leu Ile Leu Gly Gly Ile Pro Asn Glu Lys Tyr
210 215 220
Val Leu Met Thr Asp Thr Ser Phe Lys Asp Phe Ser Leu Val Glu Val
225 230 235 240
Asn Gly Val Gly Gln Met Leu Ser Ile Asp Ser Cys Trp Val Gly Ser
245 250 255
Phe Tyr Cys Pro His Ser Gly Phe Thr Ala Thr Ile Tyr Asp Thr Ile
260 265 270
Ala Thr Glu Ser Thr Leu Phe Ile Arg Gln Asn Gln Leu Val Tyr Tyr
275 280 285
Phe Thr Gly Thr Tyr Thr Thr Leu Tyr Glu Arg Asn Arg Gly Ser Gly
290 295 300
Glu Cys Ala Val Ala Gly Pro Thr Pro Gly Glu Gly Thr Leu Val Asn
305 310 315 320
Pro Ser Thr Glu Gly Ser Trp Ile Arg Val Leu Ala Ser Glu Cys Ile
325 330 335
Lys Lys Leu Cys Pro Val Tyr Phe His Ser Asn Gly Ser Glu Tyr Ile
340 345 350
Met Ala Leu Thr Thr Gly Lys His Glu Gly Tyr Val His Phe Gly Thr
355 360 365
Ile Arg Val Thr Thr Cys Ser Ile Ile Trp Ser Glu Tyr Ile Ala Gly
370 375 380
Glu Tyr Thr Leu Leu Leu Leu Val Glu Ser Gly Tyr Gly Asn Ala Ser

385 390 395 400
Lys Arg Phe Gln Val Val Ser Tyr Asn Thr Ala Ser Asp Asp Leu Glu
405 410 415
Leu Leu Tyr His Ile Pro Glu Phe Ile Pro Glu Ala Arg Gly Leu Glu
420 425 430
Phe Leu Met Ile Leu Gly Thr Glu Ser Tyr Thr Ser Thr Ala Met Ala
435 440 445
Pro Lys Gly Ile Phe Cys Asn Pro Tyr Asn Asn Leu Ile Phe Ile Trp
450 455 460
Gly Asn Phe Leu Leu Gln Ser Ser Asn Lys Glu Asn Phe Ile Tyr Leu
465 470 475 480
Ala Asp Phe Pro Lys Glu Leu Ser Ile Lys Tyr Met Ala Arg Ser Phe
485 490 495
Arg Gly Ala Val Ala Ile Val Thr Glu Thr Glu Glu Ile Trp Tyr Leu
500 505 510
Leu Glu Gly Ser Tyr Arg Val Tyr Gln Leu Phe Pro Ser Lys Gly Trp
515 520 525
Gln Val His Ile Ser Leu Lys Leu Met Gln Gln Ser Ser Leu Tyr Ala
530 535 540
Ser Asn Glu Thr Met Leu Thr Leu Phe Tyr Glu Asp Ser Lys Leu Tyr
545 550 555 560
Gln Leu Val Tyr Leu Met Asn Asn Gln Lys Gly Gln Leu Val Lys Arg
565 570 575
Leu Val Pro Val Glu Gln Leu Leu Met Tyr Gln Gln His Thr Ser His
580 585 590
Tyr Asp Leu Glu Arg Lys Gly Gly Tyr Leu Met Leu Ser Phe Ile Asp
595 600 605
Phe Cys Pro Phe Ser Val Met Arg Leu Arg Ser Leu Pro Ser Pro Gln
610 615 620
Arg Tyr Thr Arg Gln Glu Arg Tyr Arg Ala Arg Pro Pro Arg Val Leu
625 630 635 640
Glu Arg Ser Gly Phe Pro Gln Gly Glu Leu Ala Arg His Leu Pro Gly
645 650 655
Pro Gly Leu Leu Pro Ala Val Ala Ala Leu Arg Val Arg Gln Ala Val
660 665 670
Arg Gly Pro Gly Ala Arg Pro His Leu Ala Leu Val Gly Glu Gln Gln
675 680 685
Thr Arg Pro Gly Leu Leu Leu Leu Leu Gly Glu Gln Leu Ala Lys Arg
690 695 700
Gly Arg Arg Val His Arg Asn Gly Gln Leu Arg Lys Asp Leu Gln Pro
705 710 715 720
Arg Val Arg Val Arg Ala Ala Gly Ala His Phe Pro Gly Gln Gly His
725 730 735 736
*

<210> 1248
<211> 175
<212> PRT
<213> Homo sapiens

<400> 1248
Met Gly Trp Val Trp Thr Leu Cys Thr Ala Ser Ala Cys Leu Thr Leu
1 5 10 15
Leu Phe Trp Ser Gln Thr Pro Gly Lys Ala Phe Gln Ile Pro Cys Pro
20 25 30

Pro Pro His Leu Ser His Trp Cys Leu Ser Pro Met Gln Met Asp Asp
35 40 45
Gly Cys Ala Arg Leu Cys Val Leu Trp Thr Ala Trp Met Arg Trp Arg
50 55 60
Val Leu Met Cys Ser Cys Arg Val Trp Ala Thr Asp Leu Gly Ile Phe
65 70 75 80
Leu Gly Val Ala Leu Gly Asn Glu Pro Leu Glu Met Trp Pro Leu Thr
85 90 95
Gln Asn Glu Glu Cys Thr Val Thr Gly Phe Leu Arg Asp Lys Leu Gln
100 105 110
Tyr Arg Ser Arg Leu Gln Tyr Met Lys His Tyr Phe Pro Ile Asn Tyr
115 120 125
Lys Ile Arg Val Pro Tyr Glu Gly Val Phe Arg Ile Ala Asn Val Thr
130 135 140
Arg Leu Arg Ala Gln Gly Ser Glu Arg Glu Leu Arg Tyr Leu Gly Val
145 150 155 160
Leu Val Ser Leu Ser Ala Thr Glu Ser Val His Asp Glu Leu Leu
165 170 175

<210> 1249
<211> 68
<212> PRT
<213> Homo sapiens

<400> 1249
Met Phe His Arg Cys Arg Leu Lys Ala Gly Leu Met Leu Trp Arg Ser
1 5 10 15
Leu Glu Ser Gly Leu Cys Ala Gly Ala His Arg Leu Trp Leu Glu Gly
20 25 30
Pro Met Ala Phe Pro Glu Leu Gly Glu Lys Asp Pro Leu Leu Ala Ser
35 40 45
Pro Leu Ala Leu Ile Pro Gln Ser Leu Ile Gly Leu Gly Gly Leu Arg
50 55 60
Gly Ala Trp *
65 67

<210> 1250
<211> 209
<212> PRT
<213> Homo sapiens

<400> 1250
Met Ser Phe Cys Phe Thr Phe Leu Ser Leu Leu Pro Ala Cys Ile Lys
1 5 10 15
Leu Ile Leu Gln Pro Ser Ser Lys Gly Phe Lys Phe Thr Leu Val Ser
20 25 30
Cys Ala Leu Ser Phe Phe Leu Phe Ser Phe Gln Val His Glu Lys Ser
35 40 45
Ile Leu Leu Val Ser Leu Pro Val Cys Leu Val Leu Ser Glu Ile Pro
50 55 60
Phe Met Ser Thr Trp Phe Leu Leu Val Ser Thr Phe Ser Met Leu Pro
65 70 75 80
Leu Leu Leu Lys Asp Glu Leu Leu Met Pro Ser Val Val Thr Thr Met

85 90 95
Ala Phe Phe Ile Ala Cys Val Thr Ser Phe Ser Ile Phe Glu Lys Thr
100 105 110
Ser Glu Glu Glu Leu Gln Leu Lys Ser Phe Ser Ile Ser Val Arg Lys
115 120 125
Tyr Leu Pro Cys Phe Thr Phe Leu Ser Arg Ile Ile Gln Tyr Leu Phe
130 135 140
Leu Ile Ser Val Ile Thr Met Val Leu Leu Thr Leu Met Thr Val Thr
145 150 155 160
Leu Asp Pro Pro Gln Lys Leu Pro Asp Leu Phe Ser Val Leu Val Cys
165 170 175
Phe Val Ser Cys Leu Asn Phe Leu Phe Phe Leu Val Tyr Phe Asn Ile
180 185 190
Ile Ile Met Trp Asp Ser Lys Ser Gly Arg Asn Gln Lys Lys Ile Ser
195 200 205 208
*

<210> 1251
<211> 58
<212> PRT
<213> Homo sapiens

<400> 1251
Met Ile Leu Leu Leu Ser Thr Phe Phe Cys Cys Phe Arg Glu Asp Ser
1 5 10 15
Cys Phe Tyr Lys Lys Tyr Val Gly Leu Val Gln Trp Leu Met Pro Val
20 25 30
Ile Pro Ala Leu Trp Glu Ala Lys Val Gly Gly Ser Leu Glu Val Trp
35 40 45
Ser Ser Arg Pro Ala Trp Pro Ile Arg *
50 55 57

<210> 1252
<211> 84
<212> PRT
<213> Homo sapiens

<400> 1252
Met Tyr Lys Asn Phe Cys Leu Phe Phe Ile Phe Ala Leu Tyr Gln Gly
1 5 10 15
Leu Ala Asn Tyr Gly Leu Trp Ala Asn Ser Asn Pro Leu His Val Ser
20 25 30
Val Tyr Lys Ile Leu Leu Gly Cys Val Pro Trp Leu Leu Ser Val Val
35 40 45
Ser Ala Ser Arg Val Ala Gly Thr Thr Gly Thr His His Tyr Ala Trp
50 55 60
Ile Ile Phe Cys Ile Phe Ser Thr Asp Gly Val Ser Pro Arg Trp Pro
65 70 75 80
Arg Trp Ser *
83

<210> 1253
<211> 73
<212> PRT
<213> Homo sapiens

<400> 1253
Met Glu Phe Gly Leu Ser Trp Leu Phe Leu Val Ala Ile Leu Lys Gly
1 5 10 15
Val Gln Cys Glu Val Gln Leu Val Glu Ser Gly Gly Gly Leu Val Gln
20 25 30
Pro Gly Gly Ser Leu Arg Leu Ser Cys Ala Ala Ser Gly Phe Thr Phe
35 40 45
Ser Ser Tyr Ala Met Ser Trp Val Arg Gln Ala Pro Gly Lys Gly Glu
50 55 60
Gly Ala Gly Val Gly Leu Arg Phe *
65 70 72

<210> 1254
<211> 209
<212> PRT
<213> Homo sapiens

<400> 1254
Met Ser Phe Cys Phe Thr Phe Leu Ser Leu Leu Pro Ala Cys Ile Lys
1 5 10 15
Leu Ile Leu Gln Pro Ser Ser Lys Gly Phe Lys Phe Thr Leu Val Ser
20 25 30
Cys Ala Leu Ser Phe Phe Leu Phe Ser Phe Gln Val His Glu Lys Ser
35 40 45
Ile Leu Leu Val Ser Leu Pro Val Cys Leu Val Leu Ser Glu Ile Pro
50 55 60
Phe Met Ser Thr Trp Phe Leu Leu Val Ser Thr Phe Ser Met Leu Pro
65 70 75 80
Leu Leu Leu Lys Asp Glu Leu Leu Met Pro Ser Val Val Thr Thr Met
85 90 95
Ala Phe Phe Ile Ala Cys Val Thr Ser Phe Ser Ile Phe Glu Lys Thr
100 105 110
Ser Glu Glu Glu Leu Gln Leu Lys Ser Phe Ser Ile Ser Val Arg Lys
115 120 125
Tyr Leu Pro Cys Phe Thr Phe Leu Ser Arg Ile Ile Gln Tyr Leu Phe
130 135 140
Leu Ile Ser Val Ile Thr Met Val Leu Leu Thr Leu Met Thr Val Thr
145 150 155 160
Leu Asp Pro Pro Gln Lys Leu Pro Asp Leu Phe Ser Val Leu Val Cys
165 170 175
Phe Val Ser Cys Leu Asn Phe Leu Phe Phe Leu Val Tyr Phe Asn Ile
180 185 190
Ile Ile Met Trp Asp Ser Lys Ser Gly Arg Asn Gln Lys Lys Ile Ser
195 200 205 208
*

<210> 1255
<211> 730
<212> PRT
<213> Homo sapiens

<400> 1255
Met Gly Pro Trp Gly Trp Lys Leu Arg Trp Thr Val Ala Leu Leu Leu
1 5 10 15
Ala Ala Ala Gly Thr Ala Val Gly Asp Arg Cys Glu Arg Asn Glu Phe
20 25 30
Gln Cys Gln Asp Gly Lys Cys Ile Ser Tyr Lys Trp Val Cys Asp Gly
35 40 45
Ser Ala Glu Cys Gln Asp Gly Ser Asp Glu Ser Gln Glu Thr Cys Leu
50 55 60
Ser Val Thr Cys Lys Ser Gly Asp Phe Ser Cys Gly Gly Arg Val Asn
65 70 75 80
Arg Cys Ile Pro Gln Phe Trp Arg Cys Asp Gly Gln Val Asp Cys Asp
85 90 95
Asn Gly Ser Asp Glu Gln Gly Cys Pro Pro Lys Thr Cys Ser Gln Asp
100 105 110
Glu Phe Arg Cys His Asp Gly Lys Cys Ile Ser Arg Gln Phe Val Cys
115 120 125
Asp Ser Asp Arg Asp Cys Leu Asp Gly Ser Asp Glu Ala Ser Cys Pro
130 135 140
Val Leu Thr Cys Gly Pro Ala Ser Phe Gln Cys Asn Ser Ser Thr Cys
145 150 155 160
Ile Pro Gln Leu Trp Ala Cys Asp Asn Asp Pro Asp Cys Glu Asp Gly
165 170 175
Ser Asp Glu Trp Pro Gln Arg Cys Arg Gly Leu Tyr Val Phe Gln Gly
180 185 190
Asp Ser Ser Pro Cys Ser Ala Phe Glu Phe His Cys Leu Ser Gly Glu
195 200 205
Cys Ile His Ser Ser Trp Arg Cys Asp Gly Gly Pro Asp Cys Lys Asp
210 215 220
Lys Ser Asp Glu Glu Asn Cys Ala Val Ala Thr Cys Arg Pro Asp Glu
225 230 235 240
Phe Gln Cys Ser Asp Gly Asn Cys Ile His Gly Ser Arg Gln Cys Asp
245 250 255
Arg Glu Tyr Asp Cys Lys Asp Met Ser Asp Glu Val Gly Cys Val Asn
260 265 270
Val Thr Leu Cys Glu Gly Pro Asn Lys Phe Lys Cys His Ser Gly Glu
275 280 285
Cys Ile Thr Leu Asp Lys Val Cys Asn Met Ala Arg Asp Cys Arg Asp
290 295 300
Trp Ser Asp Glu Pro Ile Lys Glu Cys Gly Thr Asn Glu Cys Leu Asp
305 310 315 320
Asn Asn Gly Gly Cys Ser His Val Cys Asn Asp Leu Lys Ile Gly Tyr
325 330 335
Glu Cys Leu Cys Pro Asp Gly Phe Gln Leu Val Ala Gln Arg Arg Cys
340 345 350
Glu Asp Ile Asp Glu Cys Gln Asp Pro Asp Thr Cys Ser Gln Leu Cys
355 360 365
Val Asn Leu Glu Gly Gly Tyr Lys Cys Gln Cys Glu Glu Gly Phe Gln
370 375 380
Leu Asp Pro His Thr Lys Ala Cys Lys Ala Val Gly Ser Ile Ala Tyr
385 390 395 400
Leu Phe Phe Thr Asn Arg His Glu Val Arg Lys Met Thr Leu Asp Arg
405 410 415

Ser Glu Tyr Thr Ser Leu Ile Pro Asn Leu Arg Asn Val Val Ala Leu
420 425 430
Asp Thr Glu Val Ala Ser Asn Arg Ile Tyr Trp Ser Asp Leu Ser Gln
435 440 445
Arg Met Ile Cys Ser Thr Gln Leu Asp Arg Ala His Gly Val Ser Ser
450 455 460
Tyr Asp Thr Val Ile Ser Arg Asp Ile Gln Ala Pro Asp Gly Leu Ala
465 470 475 480
Val Asp Trp Ile His Ser Asn Ile Tyr Trp Thr Asp Ser Val Leu Gly
485 490 495
Thr Val Ser Val Ala Asp Thr Lys Gly Val Lys Arg Lys Thr Leu Phe
500 505 510
Arg Glu Asn Gly Ser Lys Pro Arg Ala Ile Val Val Asp Pro Val His
515 520 525
Gly Phe Met Tyr Trp Thr Asp Trp Gly Thr Pro Ala Lys Ile Lys Lys
530 535 540
Gly Gly Leu Asn Gly Val Asp Ile Tyr Ser Leu Val Thr Glu Asn Ile
545 550 555 560
Gln Trp Pro Asn Gly Ile Thr Leu Asp Leu Leu Ser Gly Arg Leu Tyr
565 570 575
Trp Val Asp Ser Lys Leu His Ser Ile Ser Ser Ile Asp Val Asn Gly
580 585 590
Gly Asn Arg Lys Thr Ile Leu Glu Asp Glu Lys Arg Leu Ala His Pro
595 600 605
Phe Ser Leu Ala Val Phe Glu Asp Lys Val Phe Trp Thr Asp Ile Ile
610 615 620
Asn Glu Ala Ile Phe Ser Ala Asn Arg Leu Thr Gly Ser Asp Val Asn
625 630 635 640
Leu Leu Ala Glu Asn Leu Leu Ser Pro Glu Asp Met Val Leu Phe His
645 650 655
Asn Leu Thr Gln Pro Arg Gly Val Asn Trp Cys Glu Arg Thr Thr Leu
660 665 670
Ser Asn Gly Gly Cys Gln Tyr Leu Cys Leu Pro Ala Pro Gln Ile Asn
675 680 685
Pro His Ser Pro Lys Phe Thr Cys Ala Cys Pro Asp Gly Met Leu Leu
690 695 700
Ala Arg Gly His Glu Glu Leu Pro His Arg Gly Leu Arg Leu Gln Trp
705 710 715 720
Pro Pro Arg Arg His Pro Pro Ser Gly *
725 729

<210> 1256
<211> 264
<212> PRT
<213> Homo sapiens

<400> 1256
Met Arg Gly Asn Leu Ala Leu Val Gly Val Leu Ile Ser Leu Ala Phe
1 5 10 15
Leu Ser Leu Leu Pro Ser Gly His Pro Gln Pro Ala Gly Asp Asp Ala
20 25 30
Cys Ser Val Gln Ile Leu Val Pro Gly Leu Lys Gly Asp Ala Gly Glu
35 40 45
Lys Gly Asp Lys Gly Ala Pro Gly Arg Pro Gly Arg Val Gly Pro Thr
50 55 60
Gly Glu Lys Gly Asp Met Gly Asp Lys Gly Gln Lys Gly Ser Val Gly

```
65                  70                  75                  80
Arg His Gly Lys Ile Gly Pro Ile Gly Ser Lys Gly Glu Lys Gly Asp
                85                  90                  95
Ser Gly Asp Ile Gly Pro Pro Gly Pro Asn Gly Glu Pro Gly Leu Pro
            100                 105                 110
Cys Glu Cys Ser Gln Leu Arg Lys Ala Ile Gly Glu Met Asp Asn Gln
        115                 120                 125
Val Ser Gln Leu Thr Ser Glu Leu Lys Phe Ile Lys Asn Ala Val Ala
    130                 135                 140
Gly Val Arg Glu Thr Glu Ser Lys Ile Tyr Leu Leu Val Lys Glu Glu
145                 150                 155                 160
Lys Arg Tyr Ala Asp Ala Gln Leu Ser Cys Gln Gly Arg Gly Gly Thr
                165                 170                 175
Leu Ser Met Pro Lys Asp Glu Ala Ala Asn Gly Leu Met Ala Ala Tyr
            180                 185                 190
Leu Ala Gln Ala Gly Leu Ala Arg Val Phe Ile Gly Ile Asn Asp Leu
        195                 200                 205
Glu Lys Glu Gly Ala Phe Val Tyr Ser Asp His Ser Pro Met Arg Thr
    210                 215                 220
Phe Asn Lys Trp Arg Ser Gly Glu Pro Asn Asn Ala Tyr Asp Glu Glu
225                 230                 235                 240
Asp Cys Val Glu Met Val Ala Ser Gly Gly Trp Asn Asp Val Ala Cys
                245                 250                 255
His Thr Thr Met Tyr Phe Met  *
            260         263
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<210> 1257
<211> 407
<212> PRT
<213> Homo sapiens
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<400> 1257
Met Ser Gly Ala Pro Thr Ala Gly Ala Ala Leu Met Leu Cys Ala Ala
 1               5                  10                  15
Thr Ala Val Leu Leu Ser Ala Gln Gly Gly Pro Val Gln Ser Lys Ser
            20                  25                  30
Pro Arg Phe Ala Ser Trp Asp Glu Met Asn Val Leu Ala His Gly Leu
        35                  40                  45
Leu Gln Leu Gly Gln Gly Leu Arg Glu His Ala Glu Arg Thr Arg Ser
    50                  55                  60
Gln Leu Ser Ala Leu Glu Arg Arg Leu Ser Ala Cys Gly Ser Ala Cys
65                  70                  75                  80
Gln Gly Thr Glu Gly Ser Thr Asp Leu Pro Leu Ala Pro Glu Ser Arg
                85                  90                  95
Val Asp Pro Glu Val Leu His Ser Leu Gln Thr Gln Leu Lys Ala Gln
            100                 105                 110
Asn Ser Arg Ile Gln Gln Leu Phe His Lys Val Ala Gln Gln Gln Arg
        115                 120                 125
His Leu Glu Lys Gln His Leu Arg Ile Gln His Leu Gln Ser Gln Phe
    130                 135                 140
Gly Leu Leu Asp His Lys His Leu Asp His Glu Val Ala Lys Pro Ala
145                 150                 155                 160
Arg Arg Lys Arg Leu Pro Glu Met Ala Gln Pro Val Asp Pro Ala His
                165                 170                 175
Asn Val Ser Arg Leu His Arg Leu Pro Arg Asp Cys Gln Glu Leu Phe
            180                 185                 190
```

Gln Val Gly Glu Arg Gln Ser Gly Leu Phe Glu Ile Gln Pro Gln Gly
195 200 205
Ser Pro Pro Phe Leu Val Asn Cys Lys Met Thr Ser Asp Gly Gly Trp
210 215 220
Thr Val Ile Gln Arg Arg His Asp Gly Ser Val Asp Phe Asn Arg Pro
225 230 235 240
Trp Glu Ala Tyr Lys Ala Gly Phe Gly Asp Pro His Gly Glu Phe Trp
245 250 255
Leu Gly Leu Glu Lys Val His Ser Ile Thr Gly Asp Arg Asn Ser Arg
260 265 270
Leu Ala Val Gln Leu Arg Asp Trp Asp Gly Asn Ala Glu Leu Leu Gln
275 280 285
Phe Ser Val His Leu Gly Gly Glu Asp Thr Ala Tyr Ser Leu Gln Leu
290 295 300
Thr Ala Pro Val Ala Gly Gln Leu Gly Ala Thr Thr Val Pro Pro Ser
305 310 315 320
Gly Leu Ser Val Pro Phe Ser Thr Trp Asp Gln Asp His Asp Leu Arg
325 330 335
Arg Asp Lys Asn Cys Ala Lys Ser Leu Ser Gly Gly Trp Trp Phe Gly
340 345 350
Thr Cys Ser His Ser Asn Leu Asn Gly Gln Tyr Phe Arg Ser Ile Pro
355 360 365
Gln Gln Arg Gln Lys Leu Lys Lys Gly Ile Phe Trp Lys Thr Trp Arg
370 375 380
Gly Arg Tyr Tyr Pro Leu Gln Ala Thr Thr Met Leu Ile Gln Pro Met
385 390 395 400
Ala Ala Glu Ala Ala Ser *
405 406

<210> 1258
<211> 120
<212> PRT
<213> Homo sapiens

<400> 1258
Met Met Thr Pro Lys Leu Met Ile Trp Leu Leu Leu Gln Ala Lys Ser
1 5 10 15
Ser Ile Ser Met Leu Glu Lys Ser Ser Lys Cys Leu Gly Arg Cys Phe
20 25 30
Ser Ser Phe Ala Lys Asn Leu Val Met Ile Gln Ser Cys Val Ser Trp
35 40 45
Ala Leu Met Ser Glu Asn Phe Tyr Arg Thr Leu Met Leu Cys Thr Thr
50 55 60
Thr Leu Leu Pro Ser Thr Gln Glu Cys Val His Leu Pro Leu Gly Ala
65 70 75 80
Leu Met Gln Lys Arg Ala Lys Asp Ser Phe Cys Thr Thr Thr Gln Arg
85 90 95
Glu Lys Asp Phe Arg Ile Leu Ser Leu Glu Ser Ser Lys Gln Trp His
100 105 110
Asn Lys Ser Met Ala Leu Lys *
115 119

<210> 1259
<211> 160

<212> PRT
<213> Homo sapiens

<400> 1259
Met Val Cys Leu Arg Leu Pro Gly Gly Ser Cys Met Ala Val Leu Thr
1 5 10 15
Val Thr Leu Met Val Leu Ser Ser Pro Leu Ala Leu Ala Gly Asp Thr
20 25 30
Arg Pro Arg Phe Leu Glu Tyr Ser Thr Gly Glu Cys Tyr Phe Phe Asn
35 40 45
Gly Thr Glu Arg Val Arg Phe Leu Asp Arg Tyr Phe Tyr Asn Gln Glu
50 55 60
Glu Tyr Val Arg Phe Asp Ser Asp Val Gly Glu Tyr Arg Ala Val Thr
65 70 75 80
Glu Leu Gly Arg Pro Asp Ala Glu Tyr Leu Glu Gln Pro Glu Gly Arg
85 90 95
Pro Trp Asn Ser Gln Lys Asp Ile Leu Glu Asp Glu Arg Ala Ala Val
100 105 110
Asp Thr Tyr Cys Arg His Asn Tyr Gly Val Val Glu Ser Phe Thr Val
115 120 125
Gln Arg Arg Val His Pro Lys Val Thr Val Tyr Pro Ser Lys Thr Gln
130 135 140
Pro Leu Gln Ala Pro Gln Pro Ala Val Leu Phe Cys Glu Trp Phe *
145 150 155 159

<210> 1260
<211> 111
<212> PRT
<213> Homo sapiens

<400> 1260
Met Leu Thr Phe Leu Met Leu Val Arg Leu Ser Thr Leu Cys Pro Ser
1 5 10 15
Ala Val Leu Gln Arg Leu Asp Arg Leu Val Glu Pro Leu Arg Ala Thr
20 25 30
Cys Thr Thr Lys Val Lys Ala Asn Ser Val Lys Gln Glu Phe Glu Lys
35 40 45
Gln Asp Glu Leu Lys Arg Ser Ala Met Arg Ala Val Ala Ala Leu Leu
50 55 60
Thr Ile Pro Glu Ala Glu Lys Ser Pro Leu Met Ser Glu Phe Gln Ser
65 70 75 80
Gln Ile Ser Ser Asn Pro Glu Leu Ala Ala Ile Phe Glu Ser Ile Gln
85 90 95
Lys Asp Ser Ser Ser Thr Asn Leu Glu Ser Met Asp Thr Ser *
100 105 110

<210> 1261
<211> 123
<212> PRT
<213> Homo sapiens

<400> 1261

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Met Ile Pro Ala Arg Phe Ala Gly Val Leu Leu Ala Leu Ala Leu Ile
1               5                   10                  15
Leu Pro Gly Thr Leu Cys Ala Glu Gly Thr Arg Gly Arg Ser Ser Thr
            20                  25                  30
Ala Arg Cys Ser Leu Phe Gly Ser Asp Phe Val Asn Thr Phe Asp Gly
        35                  40                  45
Ser Met Tyr Ser Phe Ala Gly Tyr Cys Ser Tyr Leu Leu Ala Gly Gly
    50                  55                  60
Cys Gln Lys Arg Ser Phe Ser Ile Ile Gly Asp Phe Gln Asn Gly Lys
65                  70                  75                  80
Arg Val Ser Leu Ser Val Tyr Leu Gly Glu Phe Phe Asp Ile His Leu
                85                  90                  95
Phe Val Asn Gly Thr Val Thr Gln Gly Asp Gln Arg Val Ser Met Pro
            100                 105                 110
Tyr Ala Ser Lys Gly Leu Tyr Leu Glu Thr  *
        115                 120     122


<210> 1262
<211> 737
<212> PRT
<213> Homo sapiens


<400> 1262
Met Phe Pro Ala Gly Pro Pro Trp Pro Arg Val Arg Val Val Gln Val
1               5                   10                  15
Leu Trp Ala Leu Leu Ala Val Leu Leu Ala Ser Trp Arg Leu Trp Ala
            20                  25                  30
Ile Lys Asp Phe Gln Glu Cys Thr Trp Gln Val Val Leu Asn Glu Phe
        35                  40                  45
Lys Arg Val Gly Glu Ser Gly Val Ser Asp Ser Phe Phe Glu Gln Glu
    50                  55                  60
Pro Val Asp Thr Val Ser Ser Leu Phe His Met Leu Val Asp Ser Pro
65                  70                  75                  80
Ile Asp Pro Ser Glu Lys Tyr Leu Gly Phe Pro Tyr Tyr Leu Lys Ile
                85                  90                  95
Asn Tyr Ser Cys Glu Glu Lys Pro Ser Glu Asp Leu Val Arg Met Gly
            100                 105                 110
His Leu Thr Gly Leu Lys Pro Leu Val Leu Val Thr Phe Gln Ser Pro
        115                 120                 125
Val Asn Phe Tyr Arg Trp Lys Ile Glu Gln Leu Gln Ile Gln Met Glu
    130                 135                 140
Ala Ala Pro Phe Arg Ser Lys Gly Gly Pro Gly Gly Gly Gly Arg Asp
145                 150                 155                 160
Arg Asn Leu Ala Gly Met Asn Ile Asn Gly Phe Leu Lys Arg Asp Arg
                165                 170                 175
Asp Asn Asn Ile Gln Phe Thr Val Gly Glu Glu Leu Phe Asn Leu Met
            180                 185                 190
Pro Gln Tyr Phe Val Gly Val Ser Ser Arg Pro Leu Trp His Thr Val
        195                 200                 205
Asp Gln Ser Pro Val Leu Ile Leu Gly Gly Ile Pro Asn Glu Lys Tyr
    210                 215                 220
Val Leu Met Thr Asp Thr Ser Phe Lys Asp Phe Ser Leu Val Glu Val
225                 230                 235                 240
Asn Gly Val Gly Gln Met Leu Ser Ile Asp Ser Cys Trp Val Gly Ser
                245                 250                 255
Phe Tyr Cys Pro His Ser Gly Phe Thr Ala Thr Ile Tyr Asp Thr Ile
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                260                 265                 270
Ala Thr Glu Ser Thr Leu Phe Ile Arg Gln Asn Gln Leu Val Tyr Tyr
            275                 280                 285
Phe Thr Gly Thr Tyr Thr Thr Leu Tyr Glu Arg Asn Arg Gly Ser Gly
        290                 295                 300
Glu Cys Ala Val Ala Gly Pro Thr Pro Gly Glu Gly Thr Leu Val Asn
305                 310                 315                 320
Pro Ser Thr Glu Gly Ser Trp Ile Arg Val Leu Ala Ser Glu Cys Ile
                    325                 330                 335
Lys Lys Leu Cys Pro Val Tyr Phe His Ser Asn Gly Ser Glu Tyr Ile
                340                 345                 350
Met Ala Leu Thr Thr Gly Lys His Glu Gly Tyr Val His Phe Gly Thr
            355                 360                 365
Ile Arg Val Thr Thr Cys Ser Ile Ile Trp Ser Glu Tyr Ile Ala Gly
        370                 375                 380
Glu Tyr Thr Leu Leu Leu Leu Val Glu Ser Gly Tyr Gly Asn Ala Ser
385                 390                 395                 400
Lys Arg Phe Gln Val Val Ser Tyr Asn Thr Ala Ser Asp Asp Leu Glu
                    405                 410                 415
Leu Leu Tyr His Ile Pro Glu Phe Ile Pro Glu Ala Arg Gly Leu Glu
                420                 425                 430
Phe Leu Met Ile Leu Gly Thr Glu Ser Tyr Thr Ser Thr Ala Met Ala
            435                 440                 445
Pro Lys Gly Ile Phe Cys Asn Pro Tyr Asn Asn Leu Ile Phe Ile Trp
        450                 455                 460
Gly Asn Phe Leu Leu Gln Ser Ser Asn Lys Glu Asn Phe Ile Tyr Leu
465                 470                 475                 480
Ala Asp Phe Pro Lys Glu Leu Ser Ile Lys Tyr Met Ala Arg Ser Phe
                    485                 490                 495
Arg Gly Ala Val Ala Ile Val Thr Glu Thr Glu Glu Ile Trp Tyr Leu
                500                 505                 510
Leu Glu Gly Ser Tyr Arg Val Tyr Gln Leu Phe Pro Ser Lys Gly Trp
            515                 520                 525
Gln Val His Ile Ser Leu Lys Leu Met Gln Gln Ser Ser Leu Tyr Ala
        530                 535                 540
Ser Asn Glu Thr Met Leu Thr Leu Phe Tyr Glu Asp Ser Lys Leu Tyr
545                 550                 555                 560
Gln Leu Val Tyr Leu Met Asn Asn Gln Lys Gly Gln Leu Val Lys Arg
                    565                 570                 575
Leu Val Pro Val Glu Gln Leu Leu Met Tyr Gln Gln His Thr Ser His
                580                 585                 590
Tyr Asp Leu Glu Arg Lys Gly Gly Tyr Leu Met Leu Ser Phe Ile Asp
            595                 600                 605
Phe Cys Pro Phe Ser Val Met Arg Leu Arg Ser Leu Pro Ser Pro Gln
        610                 615                 620
Arg Tyr Thr Arg Gln Glu Arg Tyr Arg Ala Arg Pro Pro Arg Val Leu
625                 630                 635                 640
Glu Arg Ser Gly Phe Pro Gln Gly Glu Leu Ala Arg His Leu Pro Gly
                    645                 650                 655
Pro Gly Leu Leu Pro Ala Val Ala Ala Leu Arg Val Arg Gln Ala Val
                660                 665                 670
Arg Gly Pro Gly Ala Arg Pro His Leu Ala Leu Val Gly Glu Gln Gln
            675                 680                 685
Thr Arg Pro Gly Leu Leu Leu Leu Leu Gly Glu Gln Leu Ala Lys Arg
        690                 695                 700
Gly Arg Arg Val His Arg Asn Gly Gln Leu Arg Lys Asp Leu Gln Pro
705                 710                 715                 720
Arg Val Arg Val Arg Ala Ala Gly Ala His Phe Pro Gly Gln Gly His
                    725                 730                 735 736
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<210> 1263
<211> 48
<212> PRT
<213> Homo sapiens

<400> 1263
Met Gly Ala Gly Cys Thr Pro Val Val Leu Gly Ala Ala Leu Trp Leu
1 5 10 15
Trp Arg Trp Phe Ser Arg Trp Gly Leu Gly Gly Leu Cys Trp Arg Pro
20 25 30
Cys Thr Cys Thr Pro Cys His Ser Ala Ser Pro Gly Ala Gly Arg *
35 40 45 47

<210> 1264
<211> 61
<212> PRT
<213> Homo sapiens

<400> 1264
Met Met Tyr Ile Leu Phe Leu Gln Ala Phe Ile Leu Asp Tyr Tyr Gln
1 5 10 15
Tyr Phe Leu Gly Leu Asn Cys Val Tyr Ser Tyr Gln Ser Lys Lys Asp
20 25 30
Phe Ser Gln Ile Trp Ser Gln Gly Trp Phe Ala Leu Leu Trp Ile Leu
35 40 45
Cys Leu Ser Arg Ile Leu Glu Ser Phe Phe Phe Leu *
50 55 60

<210> 1265
<211> 58
<212> PRT
<213> Homo sapiens

<400> 1265
Met Val Gly Phe Leu ~~Cys Cys Phe Tyr~~ Leu Phe Gln Leu Leu Gly Pro
1 5 10 15
Gly Leu Leu Cys Leu Pro Lys Ala Val Leu Ser Phe Leu Gly Leu Leu
20 25 30
Glu Ala Ala His His Leu Leu Val Lys Gly Phe Leu Leu Pro Val Leu
35 40 45
Asp Leu Pro Gln Val Ile Val His Gln *
50 55 57

<210> 1266
<211> 148

<212> PRT
<213> Homo sapiens

<400> 1266
Met Ala Leu Gln Leu Trp Ala Leu Thr Leu Leu Gly Leu Leu Gly Ala
1 5 10 15
Gly Ala Ser Leu Arg Pro Arg Lys Leu Asp Phe Phe Arg Ser Glu Lys
20 25 30
Glu Leu Asn His Leu Ala Val Asp Glu Ala Ser Gly Val Val Tyr Leu
35 40 45
Gly Ala Val Asn Ala Leu Tyr Gln Leu Asp Ala Lys Leu Gln Leu Glu
50 55 60
Gln Gln Val Ala Thr Gly Pro Val Leu Asp Asn Lys Lys Cys Thr Pro
65 70 75 80
Pro Ile Glu Ala Ser Gln Cys His Glu Ala Glu Met Thr Asp Asn Val
85 90 95
Asn Gln Leu Leu Leu Val Asp Pro Pro Arg Lys Arg Leu Val Glu Cys
100 105 110
Gly Gln Leu Leu Lys Gly Ile Leu Arg Ser Ala Arg Pro Glu Gln His
115 120 125
Leu Pro Pro Pro Val Leu Arg Gly Arg Gln Arg Gly Glu Val Phe Arg
130 135 140
Gly Gln Gln *
145 147

<210> 1267
<211> 227
<212> PRT
<213> Homo sapiens

<400> 1267
Met Arg Trp Leu Trp Pro Leu Ala Val Ser Leu Ala Val Ile Leu Ala
1 5 10 15
Val Gly Leu Ser Arg Val Ser Gly Gly Ala Pro Leu His Leu Gly Arg
20 25 30
His Arg Ala Glu Thr Gln Glu Gln Gln Ser Arg Ser Lys Arg Gly Thr
35 40 45
Glu Asp Glu Glu Ala Lys Gly Val Gln Gln Tyr Val Pro Glu Glu Trp
50 55 60
Ala Glu Tyr Pro Arg Pro Ile His Pro Ala Gly Leu Gln Pro Thr Lys
65 70 75 80
Pro Leu Val Ala Thr Ser Pro Asn Pro Asp Lys Asp Gly Gly Thr Pro
85 90 95
Asp Ser Gly Gln Glu Leu Arg Gly Asn Leu Thr Gly Ala Pro Gly Gln
100 105 110
Arg Leu Gln Ile Gln Asn Pro Leu Tyr Pro Val Thr Glu Ser Ser Tyr
115 120 125
Ser Ala Tyr Ala Ile Met Leu Leu Ala Leu Val Glu Phe Ala Ala Gly
130 135 140
Ile Val Gly Asn Leu Ser Val Met Cys Ile Ala Trp His Ser Tyr Tyr
145 150 155 160
Leu Lys Ser Ala Trp Asn Ser Ile Leu Ala Ser Leu Ala Leu Trp Asp
165 170 175
Phe Leu Val Leu Phe Phe Cys Leu Pro Ile Val Ile Leu Asn Glu Ile
180 185 190

Thr Lys Gln Arg Leu Leu Gly Asp Ala Pro Cys Pro Cys Arg Ala Leu
195 200 205
His Gly Gly Leu Leu Ser Gly Ser His Asp Phe Gln Pro Leu Cys Pro
210 215 220
Gly His *
225 226

<210> 1268
<211> 983
<212> PRT
<213> Homo sapiens

<400> 1268
Met Leu Gly Asn Val Leu Leu Leu Cys Phe Phe Val Phe Phe Ile Phe
1 5 10 15
Gly Ile Val Gly Val Gln Leu Trp Ala Gly Leu Leu Arg Asn Arg Cys
20 25 30
Phe Leu Pro Glu Asn Phe Ser Leu Pro Leu Ser Val Asp Leu Glu Arg
35 40 45
Tyr Tyr Gln Thr Glu Asn Glu Asp Glu Ser Pro Phe Ile Cys Ser Gln
50 55 60
Pro Arg Glu Asn Gly Met Arg Ser Cys Arg Ser Val Pro Thr Leu Arg
65 70 75 80
Gly Asp Gly Gly Gly Gly Pro Pro Cys Gly Leu Asp Tyr Glu Ala Tyr
85 90 95
Asn Ser Ser Ser Asn Thr Thr Cys Val Asn Trp Asn Gln Tyr Tyr Thr
100 105 110
Asn Cys Ser Ala Gly Glu His Asn Pro Phe Lys Gly Ala Ile Asn Phe
115 120 125
Asp Asn Ile Gly Tyr Ala Trp Ile Ala Ile Phe Gln Val Ile Thr Leu
130 135 140
Glu Gly Trp Val Asp Ile Met Tyr Phe Val Met Asp Ala His Ser Phe
145 150 155 160
Tyr Asn Phe Ile Tyr Phe Ile Leu Leu Ile Ile Val Gly Ser Phe Phe
165 170 175
Met Ile Asn Leu Cys Leu Val Val Ile Ala Thr Gln Phe Ser Glu Thr
180 185 190
Lys Gln Arg Glu Ser Gln Leu Met Arg Glu Gln Arg Val Arg Phe Leu
195 200 205
Ser Asn Ala Ser Thr Leu Ala Ser Phe Ser Glu Pro Gly Ser Cys Tyr
210 215 220
Glu Glu Leu Leu Lys Tyr Leu Val Tyr Ile Leu Arg Lys Ala Ala Arg
225 230 235 240
Arg Leu Ala Gln Val Ser Arg Ala Ala Gly Val Arg Val Gly Leu Leu
245 250 255
Ser Ser Pro Ala Pro Leu Gly Gly Gln Glu Thr Gln Pro Ser Ser Ser
260 265 270
Cys Ser Arg Ser His Arg Arg Leu Ser Val His His Leu Val His His
275 280 285
His His His His His His His Tyr His Leu Gly Asn Gly Thr Leu Arg
290 295 300
Ala Pro Arg Ala Ser Pro Glu Ile Gln Asp Arg Asp Ala Asn Gly Ser
305 310 315 320
Arg Arg Leu Met Leu Pro Pro Pro Ser Thr Pro Ala Leu Ser Gly Ala
325 330 335
Pro Pro Gly Gly Ala Glu Ser Val His Ser Phe Tyr His Ala Asp Cys

340 345 350
His Leu Glu Pro Val Arg Cys Gln Ala Pro Pro Pro Arg Ser Pro Ser
355 360 365
Glu Ala Ser Gly Arg Thr Val Gly Ser Gly Lys Val Tyr Pro Thr Val
370 375 380
His Thr Ser Pro Pro Pro Glu Thr Leu Lys Glu Lys Ala Leu Val Glu
385 390 395 400
Val Ala Ala Ser Ser Gly Pro Pro Thr Leu Thr Ser Leu Asn Ile Pro
405 410 415
Pro Gly Pro Tyr Ser Ser Met His Lys Leu Leu Glu Thr Gln Ser Thr
420 425 430
Gly Ala Cys Gln Ser Ser Cys Lys Ile Ser Ser Pro Cys Leu Lys Ala
435 440 445
Asp Ser Gly Ala Cys Gly Pro Asp Ser Cys Pro Tyr Cys Ala Arg Ala
450 455 460
Gly Ala Gly Glu Val Glu Leu Ala Asp Arg Glu Met Pro Asp Ser Asp
465 470 475 480
Ser Glu Ala Val Tyr Glu Phe Thr Gln Asp Ala Gln His Ser Asp Leu
485 490 495
Arg Asp Pro His Ser Arg Arg Gln Arg Ser Leu Gly Pro Asp Ala Glu
500 505 510
Pro Ser Ser Val Leu Ala Phe Trp Arg Leu Ile Cys Asp Thr Phe Arg
515 520 525
Lys Ile Val Asp Ser Lys Tyr Phe Gly Arg Gly Ile Met Ile Ala Ile
530 535 540
Leu Val Asn Thr Leu Ser Met Gly Ile Glu Tyr His Glu Gln Pro Glu
545 550 555 560
Glu Leu Thr Asn Ala Leu Glu Ile Ser Asn Ile Val Phe Thr Ser Leu
565 570 575
Phe Ala Leu Glu Met Leu Leu Lys Leu Leu Val Tyr Gly Pro Phe Gly
580 585 590
Tyr Ile Lys Asn Pro Tyr Asn Ile Phe Asp Gly Val Ile Val Val Ile
595 600 605
Ser Val Trp Glu Ile Val Gly Gln Gln Gly Gly Gly Leu Ser Val Leu
610 615 620
Arg Thr Phe Arg Leu Met Arg Val Leu Lys Leu Val Arg Phe Leu Pro
625 630 635 640
Ala Leu Gln Arg Gln Leu Val Val Leu Met Lys Thr Met Asp Asn Val
645 650 655
Ala Thr Phe Cys Met Leu Leu Met Leu Phe Ile Phe Ile Phe Ser Ile
660 665 670
Leu Gly Met His Leu Phe Gly Cys Lys Phe Ala Ser Glu Arg Asp Gly
675 680 685
Asp Thr Leu Pro Asp Arg Lys Asn Phe Asp Ser Leu Leu Trp Ala Ile
690 695 700
Val Thr Val Phe Gln Ile Leu Thr Gln Glu Asp Trp Asn Lys Val Leu
705 710 715 720
Tyr Asn Gly Met Ala Ser Thr Ser Ser Trp Ala Ala Leu Tyr Phe Ile
725 730 735
Ala Leu Met Thr Phe Gly Asn Tyr Val Leu Phe Asn Leu Leu Val Ala
740 745 750
Ile Leu Val Glu Gly Phe Gln Ala Glu Gly Asp Ala Asn Lys Ser Glu
755 760 765
Ser Glu Pro Asp Phe Phe Ser Pro Ser Leu Asp Gly Asp Gly Asp Arg
770 775 780
Lys Lys Cys Leu Ala Leu Val Ser Leu Gly Glu His Pro Glu Leu Arg
785 790 795 800
Lys Ser Leu Leu Pro Pro Leu Ile Ile His Thr Ala Ala Thr Pro Met
805 810 815

Ser Leu Pro Lys Ser Thr Ser Thr Gly Leu Gly Glu Ala Leu Gly Pro
820 825 830
Ala Ser Arg Arg Thr Ser Ser Ser Gly Ser Ala Glu Pro Gly Ala Ala
835 840 845
His Glu Met Lys Ser Pro Pro Ser Ala Arg Ser Ser Pro His Ser Pro
850 855 860
Trp Ser Ala Ala Ser Ser Trp Thr Ser Arg Arg Ser Ser Arg Asn Ser
865 870 875 880
Leu Gly Arg Ala Pro Ser Leu Lys Arg Arg Ser Pro Ser Gly Glu Arg
885 890 895
Arg Ser Leu Leu Ser Gly Glu Gly Gln Glu Ser Gln Asp Glu Glu Glu
900 905 910
Ser Ser Glu Glu Glu Arg Ala Ser Pro Ala Gly Ser Asp His Arg His
915 920 925
Arg Gly Ser Leu Glu Arg Glu Ala Lys Ser Ser Phe Asp Leu Pro Asp
930 935 940
Thr Leu Gln Val Pro Gly Leu His Arg Thr Ala Ser Gly Arg Gly Ser
945 950 955 960
Ala Ser Glu His Gln Gly Leu Gln Trp Gln Val Gly Phe Arg Ala Pro
965 970 975
Gly Pro Gly Pro Ala Ala *
980 982

<210> 1269
<211> 708
<212> PRT
<213> Homo sapiens

<400> 1269
Met Leu Ser Leu Arg Arg Cys Thr Ser Met Arg Leu Cys Leu Ser Ser
1 5 10 15
Ser Leu Ala Ser Pro Cys Ser Thr Met Leu Ser Thr Val Val Leu Tyr
20 25 30
Lys Val Cys Asn Ser Phe Val Glu Met Gly Ser Ala Asn Val Gln Ala
35 40 45
Thr Asp Tyr Leu Lys Gly Val Ala Ser Leu Phe Val Val Ser Leu Gly
50 55 60
Gly Ala Ala Val Gly Leu Val Phe Ala Phe Leu Leu Ala Leu Thr Thr
65 70 75 80
Arg Phe Thr Lys Arg Val Arg Ile Ile Glu Pro Leu Leu Val Phe Leu
85 90 95
Leu Ala Tyr Ala Ala Tyr Leu Thr Ala Glu Met Ala Ser Leu Ser Ala
100 105 110
Ile Leu Ala Val Thr Met Cys Gly Leu Gly Cys Lys Lys Tyr Val Glu
115 120 125
Ala Asn Ile Ser His Lys Ser Arg Thr Thr Val Lys Tyr Thr Met Lys
130 135 140
Thr Leu Ala Ser Cys Ala Glu Thr Val Ile Phe Met Leu Leu Gly Ile
145 150 155 160
Ser Thr Val Asp Ser Ser Lys Trp Ala Trp Asp Ser Gly Leu Val Leu
165 170 175
Gly Thr Leu Ile Phe Ile Leu Phe Phe Arg Ala Leu Gly Val Val Leu
180 185 190
Gln Thr Trp Val Leu Asn Gln Phe Arg Leu Val Pro Leu Asp Lys Ile
195 200 205
Asp Gln Val Val Met Ser Tyr Gly Gly Leu Arg Gly Ala Val Ala Phe

210 215 220
Ala Leu Val Ile Leu Leu Asp Arg Thr Lys Val Pro Ala Lys Asp Tyr
225 230 235 240
Phe Val Ala Thr Thr Ile Val Val Val Phe Phe Thr Val Ile Val Gln
245 250 255
Gly Leu Thr Ile Lys Pro Leu Val Lys Trp Leu Lys Val Lys Arg Ser
260 265 270
Glu His His Lys Pro Thr Leu Asn Gln Glu Leu His Glu His Thr Phe
275 280 285
Asp His Ile Leu Ala Ala Val Glu Asp Val Val Gly His His Gly Tyr
290 295 300
His Tyr Trp Arg Asp Arg Trp Glu Gln Phe Asp Lys Lys Tyr Leu Ser
305 310 315 320
Gln Leu Leu Met Arg Arg Ser Ala Tyr Arg Ile Arg Asp Gln Ile Trp
325 330 335
Asp Val Tyr Tyr Arg Leu Asn Ile Arg Asp Ala Ile Ser Phe Val Asp
340 345 350
Gln Gly Gly His Val Leu Ser Ser Thr Gly Leu Thr Leu Pro Ser Met
355 360 365
Pro Ser Arg Asn Ser Val Ala Glu Thr Ser Val Thr Asn Leu Leu Arg
370 375 380
Glu Ser Gly Ser Gly Ala Cys Leu Asp Leu Gln Val Ile Asp Thr Val
385 390 395 400
Arg Ser Gly Arg Asp Arg Glu Asp Ala Val Met His His Leu Leu Cys
405 410 415
Gly Gly Leu Tyr Lys Pro Arg Arg Arg Tyr Lys Ala Ser Cys Ser Arg
420 425 430
His Phe Ile Ser Glu Asp Ala Gln Glu Arg Gln Asp Lys Glu Val Phe
435 440 445
Gln Gln Asn Met Lys Arg Arg Leu Glu Ser Phe Lys Ser Thr Lys His
450 455 460
Asn Ile Cys Phe Thr Lys Ser Lys Pro Arg Pro Arg Lys Thr Gly Arg
465 470 475 480
Arg Lys Lys Asp Gly Val Ala Asn Ala Glu Ala Thr Asn Gly Lys His
485 490 495
Arg Gly Leu Gly Phe Gln Asp Thr Ala Ala Val Ile Leu Thr Val Glu
500 505 510
Ser Glu Glu Glu Glu Glu Glu Ser Asp Ser Ser Glu Thr Glu Lys Glu
515 520 525
Asp Asp Glu Gly Ile Ile Phe Val Ala Arg Ala Thr Ser Glu Val Leu
530 535 540
Gln Glu Gly Lys Val Ser Gly Ser Leu Glu Val Cys Pro Ser Pro Arg
545 550 555 560
Ile Ile Pro Pro Ser Pro Thr Cys Ala Glu Lys Glu Leu Pro Trp Lys
565 570 575
Ser Gly Gln Gly Asp Leu Ala Val Tyr Val Ser Ser Glu Thr Thr Lys
580 585 590
Ile Val Pro Val Asp Met Gln Thr Gly Trp Asn Gln Ser Ile Ser Ser
595 600 605
Leu Glu Ser Leu Ala Ser Pro Pro Cys Asn Gln Ala Pro Ile Leu Thr
610 615 620
Cys Leu Pro Pro His Pro Arg Gly Thr Glu Glu Pro Gln Val Pro Leu
625 630 635 640
His Leu Pro Ser Asp Pro Arg Ser Ser Phe Ala Phe Pro Pro Ser Leu
645 650 655
Ala Lys Ala Gly Arg Ser Arg Ser Glu Ser Ser Ala Asp Leu Pro Gln
660 665 670
Gln Gln Glu Leu Gln Pro Leu Met Gly His Lys Asp His Thr His Leu
675 680 685

Ser Pro Gly Thr Ala Thr Ser His Trp Cys Ile Gln Phe Asn Arg Gly
690 695 700
Ser Arg Leu *
705 707

<210> 1270
<211> 93
<212> PRT
<213> Homo sapiens

<400> 1270
Met Leu Gln Ala Ala Leu Trp Cys Gly Ile Gly Leu Tyr Leu Val Thr
1 5 10 15
Leu Arg Leu Gly Val Glu Val Thr Pro Glu Ser Gln His Phe Gly Arg
20 25 30
Pro Arg Arg Ala Asp His Leu Arg Pro Gly Gly Arg Gly Gln Ser Gly
35 40 45
Gln His Gly Glu Thr Pro Ser Leu Leu Glu Ile Gln Lys Ile Ser Trp
50 55 60
Met Trp Trp His Ile Pro Val Ile Pro Ala Thr Trp Glu Ala Glu Ala
65 70 75 80
Gly Glu Ser Leu Glu Arg Gly Arg Trp Arg Leu Gln *
85 90 92

<210> 1271
<211> 648
<212> PRT
<213> Homo sapiens

<400> 1271
Met Leu Trp Val Thr Gly Pro Val Leu Ala Val Ile Leu Ile Ile Leu
1 5 10 15
Ile Val Ile Ala Ile Leu Leu Phe Lys Arg Lys Arg Thr His Ser Pro
20 25 30
Ser Ser Lys Asp Glu Gln Ser Ile Gly Leu Lys Asp Ser Leu Leu Ala
35 40 45
His Ser Ser Asp Pro Val Glu Met Arg Arg Leu Asn Tyr Gln Thr Pro
50 55 60
Gly Met Arg Asp His Pro Pro Ile Pro Ile Thr Asp Leu Ala Asp Asn
65 70 75 80
Ile Glu Arg Leu Lys Ala Asn Asp Gly Leu Lys Phe Ser Gln Glu Tyr
85 90 95
Glu Ser Ile Asp Pro Gly Gln Gln Phe Thr Trp Glu Asn Ser Asn Leu
100 105 110
Glu Val Asn Lys Pro Lys Asn Arg Tyr Ala Asn Val Ile Ala Tyr Asp
115 120 125
His Ser Arg Val Ile Leu Thr Ser Ile Asp Gly Val Pro Gly Ser Asp
130 135 140
Tyr Ile Asn Ala Asn Tyr Ile Asp Gly Tyr Arg Lys Gln Asn Ala Tyr
145 150 155 160
Ile Ala Thr Gln Gly Pro Leu Pro Glu Thr Met Gly Asp Phe Trp Arg
165 170 175
Met Val Trp Glu Gln Arg Thr Ala Thr Val Val Met Met Thr Arg Leu

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            180                     185                     190
Glu Glu Lys Ser Arg Val Lys Cys Asp Gln Tyr Trp Pro Ala Arg Gly
        195                     200                     205
Thr Glu Thr Cys Gly Leu Ile Gln Val Thr Leu Leu Asp Thr Val Glu
    210                     215                     220
Leu Ala Thr Tyr Thr Val Arg Thr Phe Ala Leu His Lys Ser Gly Ser
225                     230                     235                     240
Ser Glu Lys Arg Glu Leu Arg Gln Phe Gln Phe Met Ala Trp Pro Asp
                245                     250                     255
His Gly Val Pro Glu Tyr Pro Thr Pro Ile Leu Ala Phe Leu Arg Arg
            260                     265                     270
Val Lys Ala Cys Asn Pro Leu Asp Ala Gly Pro Met Val Val His Cys
        275                     280                     285
Ser Ala Gly Val Gly Arg Thr Gly Cys Phe Ile Val Ile Asp Ala Met
    290                     295                     300
Leu Glu Arg Met Lys His Glu Lys Thr Val Asp Ile Tyr Gly His Val
305                     310                     315                     320
Thr Cys Met Arg Ser Gln Arg Asn Tyr Met Val Gln Thr Glu Asp Gln
                325                     330                     335
Tyr Val Phe Ile His Glu Ala Leu Leu Glu Ala Ala Thr Cys Gly His
            340                     345                     350
Thr Glu Val Pro Ala Arg Asn Leu Tyr Ala His Ile Gln Lys Leu Gly
        355                     360                     365
Gln Val Pro Pro Gly Glu Ser Val Thr Ala Met Glu Leu Glu Phe Lys
    370                     375                     380
Leu Leu Ala Ser Ser Lys Ala His Thr Ser Arg Phe Ile Ser Ala Asn
385                     390                     395                     400
Leu Pro Cys Asn Lys Phe Lys Asn Arg Leu Val Asn Ile Met Pro Tyr
                405                     410                     415
Glu Leu Thr Arg Val Cys Leu Gln Pro Ile Arg Gly Val Glu Gly Ser
            420                     425                     430
Asp Tyr Ile Asn Ala Ser Phe Leu Asp Gly Tyr Arg Gln Gln Lys Ala
        435                     440                     445
Tyr Ile Ala Thr Gln Gly Pro Leu Ala Glu Ser Thr Glu Asp Phe Trp
    450                     455                     460
Arg Met Leu Trp Glu His Asn Ser Thr Ile Ile Val Met Leu Thr Lys
465                     470                     475                     480
Leu Arg Glu Met Gly Arg Glu Lys Cys His Gln Tyr Trp Pro Ala Glu
                485                     490                     495
Arg Ser Ala Arg Tyr Gln Tyr Phe Val Val Asp Pro Met Ala Glu Tyr
            500                     505                     510
Asn Met Pro Gln Tyr Ile Leu Arg Glu Phe Lys Val Thr Asp Ala Arg
        515                     520                     525
Asp Gly Gln Ser Arg Thr Ile Arg Gln Phe Gln Phe Thr Asp Trp Pro
    530                     535                     540
Glu Gln Gly Val Pro Lys Thr Gly Glu Gly Phe Ile Asp Phe Ile Gly
545                     550                     555                     560
Gln Val His Lys Thr Lys Glu Gln Phe Gly Gln Asp Gly Pro Ile Thr
                565                     570                     575
Val His Cys Ser Ala Gly Val Gly Arg Thr Gly Val Phe Ile Thr Leu
            580                     585                     590
Ser Ile Val Leu Glu Arg Met Arg Tyr Glu Gly Val Val Asp Met Phe
        595                     600                     605
Gln Thr Val Lys Thr Leu Arg Thr Gln Arg Pro Ala Met Val Gln Thr
    610                     615                     620
Glu Asp Gln Tyr Gln Leu Cys Tyr Arg Ala Ala Leu Glu Tyr Leu Gly
625                     630                     635                     640
Ser Phe Asp His Tyr Ala Thr  *
                645     647
```

<210> 1272
<211> 109
<212> PRT
<213> Homo sapiens

<400> 1272
Met Lys Ala Leu Cys Leu Leu Leu Leu Pro Val Leu Gly Leu Leu Val
1 5 10 15
Ser Ser Lys Thr Leu Cys Ser Met Glu Glu Ala Ile Asn Glu Arg Ile
20 25 30
Gln Glu Val Ala Gly Ser Leu Ile Phe Arg Ala Ile Ser Ser Ile Gly
35 40 45
Leu Glu Cys Gln Ser Val Thr Ser Arg Gly Asp Leu Ala Thr Cys Pro
50 55 60
Arg Gly Phe Ala Val Thr Gly Cys Thr Cys Gly Ser Ala Cys Gly Ser
65 70 75 80
Trp Asp Val Arg Ala Glu Thr Thr Cys His Cys Gln Cys Ala Gly Met
85 90 95
Asp Trp Thr Gly Ala Arg Cys Cys Arg Val Gln Pro *
100 105 108

<210> 1273
<211> 56
<212> PRT
<213> Homo sapiens

<400> 1273
Met Phe Phe Val Pro Ile Leu Leu Cys Leu Leu Leu Leu Ile Tyr Asn
1 5 10 15
Ile Ile Cys Phe Asn Met Glu His Pro Thr Gly Ala Gly Leu Arg Cys
20 25 30
Ser Leu Leu Ala Ala Pro Lys Glu Arg Gln His Arg His His Phe Val
35 40 45
Phe His Ile Asp Thr Asn His *
50 55

<210> 1274
<211> 188
<212> PRT
<213> Homo sapiens

<400> 1274
Met Asp Leu Ser Leu Leu Trp Val Leu Leu Pro Leu Val Thr Met Ala
1 5 10 15
Trp Gly Gln Tyr Gly Asp Tyr Gly Tyr Pro Tyr Gln Gln Tyr His Asp
20 25 30
Tyr Ser Asp Asp Gly Trp Val Asn Leu Asn Arg Gln Gly Phe Ser Tyr
35 40 45
Gln Cys Pro Gln Gly Gln Val Ile Val Ala Val Arg Ser Ile Phe Ser

```
     50                  55                  60
Lys Lys Glu Gly Ser Asp Arg Gln Trp Asn Tyr Ala Cys Met Pro Thr
 65                  70                  75                  80
Pro Gln Ser Leu Gly Glu Pro Thr Glu Cys Trp Trp Glu Glu Ile Asn
                 85                  90                  95
Arg Ala Gly Met Glu Trp Tyr Gln Thr Cys Ser Asn Asn Gly Leu Val
            100                 105                 110
Ala Gly Phe Gln Ser Arg Tyr Phe Glu Ser Val Leu Asp Arg Glu Trp
        115                 120                 125
Gln Phe Tyr Cys Cys Arg Tyr Ser Lys Arg Cys Pro Tyr Ser Cys Trp
    130                 135                 140
Leu Thr Thr Glu Tyr Pro Gly His Tyr Gly Glu Glu Met Asp Met Ile
145                 150                 155                 160
Ser Tyr Asn Tyr Asp Tyr Tyr Ile Arg Gly Ala Thr Thr His Phe Leu
                165                 170                 175
Cys Ser Gly Lys Gly Ser Pro Ser Gly Ser Ser  *
            180                 185     187
```

<210> 1275
<211> 81
<212> PRT
<213> Homo sapiens

<400> 1275
```
Met Val Ala Leu Thr Ile Gln Thr Trp His Trp Leu Met Thr Val Ala
 1               5                  10                  15
Glu Leu Leu Ser Leu Ala Cys Tyr Ile Ala Ser Leu Val Phe Leu His
            20                  25                  30
Glu Phe Ile Asp Val Tyr Phe Ile Ala Thr Leu Ser Phe Leu Trp Lys
        35                  40                  45
Val Ser Val Ile Thr Leu Val Ser Cys Leu Pro Leu Tyr Val Leu Lys
    50                  55                  60
Tyr Leu Arg Arg Arg Phe Ser Pro Pro Ser Tyr Ser Lys Leu Thr Ser
65                  70                  75                  80
 *
```

<210> 1276
<211> 46
<212> PRT
<213> Homo sapiens

<400> 1276
```
Met Leu Asp Leu Val Ala Leu Leu Tyr Gln Ala Val Leu Leu Pro Ala
 1               5                  10                  15
Ile Leu Leu Leu Pro Leu Cys Gln Leu Glu Met Phe Leu Met Leu Gln
            20                  25                  30
Leu Asn Arg Gln Ser Leu Lys Lys Lys Tyr Leu Ile Leu  *
        35                  40                  45
```

<210> 1277

<211> 431
<212> PRT
<213> Homo sapiens

<400> 1277
Met Ala Leu Leu Val Pro Leu Ala Leu Leu Val Ile Gln Ala His Leu
1 5 10 15
Val Leu Ser Val Gln Leu Glu Arg Val Val Thr Glu Glu Lys Val Ala
20 25 30
Leu Leu Ala Leu Leu Val Leu Pro Val Leu Leu Val Pro Glu Val Leu
35 40 45
Leu Val Leu Lys Ala His Val Val Thr Lys Val Lys Gln Val Asn Val
50 55 60
Glu Leu Leu Ala Ser Lys Asp Ile Glu Asp Ser Leu Val Ile Gln Val
65 70 75 80
Pro Gln Val Leu Gln Ala Leu Leu Val Ser Arg Val Gln Ser Ala Val
85 90 95
Gln Asp Leu Gln Ala Pro Glu Asp Leu Leu Asp Pro Val Asp Leu Leu
100 105 110
Ala Lys Met Glu Pro Val Asp Ile Gln Val Pro Leu Asp His Gln Gly
115 120 125
Leu Glu Val Thr Glu Val Lys Glu Asp Leu Arg Ala Pro Gln Ala Thr
130 135 140
Gln Gly Asn Gln Ala Leu Leu Asp Leu Leu Val Pro Leu Val Leu Ala
145 150 155 160
Val Val Val Leu Glu Pro Leu Pro Leu Leu Gly Leu Glu Val Lys Lys
165 170 175
Leu Ala Gly Phe Ala Pro Tyr Tyr Gly Asp Glu Pro Met Asp Phe Lys
180 185 190
Ile Asn Thr Asp Glu Ile Met Thr Ser Leu Lys Ser Val Asn Gly Gln
195 200 205
Ile Glu Ser Leu Ile Ser Pro Asp Gly Ser Arg Lys Asn Pro Ala Arg
210 215 220
Asn Cys Arg Asp Leu Lys Phe Cys His Pro Glu Leu Lys Ser Gly Glu
225 230 235 240
Tyr Trp Val Asp Pro Asn Gln Gly Cys Lys Leu Asp Ala Ile Lys Val
245 250 255
Phe Cys Asn Met Glu Thr Gly Glu Thr Cys Ile Ser Ala Asn Pro Leu
260 265 270
Asn Val Pro Arg Lys His Trp Trp Thr Asp Ser Ser Ala Glu Lys Lys
275 280 285
His Val Trp Phe Gly Glu Ser Met Asp Gly Gly Phe Gln Phe Ser Tyr
290 295 300
Gly Asn Pro Glu Leu Pro Glu Asp Val Leu Asp Val Gln Leu Ala Phe
305 310 315 320
Leu Arg Leu Leu Ser Ser Arg Ala Ser Gln Asn Ile Thr Tyr His Cys
325 330 335
Lys Asn Ser Ile Ala Tyr Met Asp Gln Ala Ser Gly Asn Val Lys Lys
340 345 350
Ala Leu Lys Leu Met Gly Ser Asn Glu Gly Glu Phe Lys Ala Glu Gly
355 360 365
Asn Ser Lys Phe Thr Tyr Thr Val Leu Glu Asp Gly Cys Thr Lys His
370 375 380
Thr Gly Glu Trp Ser Lys Thr Val Phe Glu Tyr Arg Thr Arg Lys Ala
385 390 395 400
Val Arg Leu Pro Ile Val Asp Ile Ala Pro Tyr Asp Ile Gly Gly Pro
405 410 415
Asp Gln Glu Phe Gly Val Asp Val Gly Pro Val Cys Phe Leu *

420 425 430

<210> 1278
<211> 53
<212> PRT
<213> Homo sapiens

<400> 1278
Met Leu Leu Tyr Val Phe Lys Phe Leu Gly Leu Phe Gln Phe Phe His
1 5 10 15
Ser Phe Cys Thr Ala Tyr Gly Pro Pro Gly Gly Cys Gly Asp Ser Gly
20 25 30
Glu Glu Thr Ser Leu Phe Phe Glu Gln Leu Asp Pro Ala Phe Trp Leu
35 40 45
Ala Asn Cys Ser *
50 52

<210> 1279
<211> 73
<212> PRT
<213> Homo sapiens

<400> 1279
Met Leu Gly Ser Ile Cys Asn Val Met Leu Leu Met Leu Ala Ala Ser
1 5 10 15
Ile Pro Glu Ile Cys Thr Phe Gly Pro Thr Lys Leu Ala Ala Asn Cys
20 25 30
Asn Trp Met Pro Ser Arg Val Ala Arg Leu Pro Ser Val Arg Asp Thr
35 40 45
Val Arg Ser Pro Pro Ala Asp Thr Glu Ala Gly Arg Ile Ala Trp Pro
50 55 60
Thr Ser Pro Gly Cys Ser Arg Phe *
65 70 72

<210> 1280
<211> 51
<212> PRT
<213> Homo sapiens

<400> 1280
Met Leu Leu Leu Leu Glu Arg Met Ala Leu Cys Pro Val Leu Asp Val
1 5 10 15
His Thr His Leu Gly Cys Ile Ile Cys Val Phe Asp Val Ala Leu Ser
20 25 30
Arg Glu Leu Ala Leu Leu Cys Arg Lys Ser Asn Trp Trp Val Ile Asn
35 40 45
Trp Leu *
50

<210> 1281
<211> 144
<212> PRT
<213> Homo sapiens

<400> 1281
Met Lys Ser Gly Ser Gly Gly Gly Ser Pro Thr Ser Leu Trp Gly Leu
1 5 10 15
Leu Phe Leu Ser Ala Ala Leu Ser Leu Trp Pro Thr Ser Gly Glu Ile
20 25 30
Cys Gly Pro Gly Ile Asp Ile Arg Asn Asp Tyr Gln Gln Leu Lys Arg
35 40 45
Leu Glu Asn Cys Thr Val Ile Glu Gly Tyr Leu His Ile Leu Leu Ile
50 55 60
Ser Lys Ala Glu Asp Tyr Arg Ser Tyr Arg Phe Pro Lys Leu Thr Val
65 70 75 80
Ile Thr Glu Tyr Leu Leu Leu Phe Arg Val Ala Gly Leu Glu Ser Leu
85 90 95
Gly Asp Leu Phe Pro Asn Leu Thr Val Ile Arg Gly Trp Lys Leu Phe
100 105 110
Tyr Asn Tyr Ala Leu Val Ile Phe Glu Met Thr Asn Leu Lys Asp Ile
115 120 125
Gly Leu Tyr Asn Leu Arg Asn Ile Thr Arg Gly Gly His Gln Asp *
130 135 140 143

<210> 1282
<211> 267
<212> PRT
<213> Homo sapiens

<400> 1282
Met Gly Pro Pro Ser Ala Cys Pro His Arg Glu Cys Ile Pro Trp Gln
1 5 10 15
Gly Leu Leu Leu Thr Ala Ser Leu Leu Thr Phe Trp Asn Ala Pro Thr
20 25 30
Thr Ala Trp Leu Phe Ile Ala Ser Ala Pro Phe Glu Val Ala Glu Gly
35 40 45
Glu Asn Val His Leu Ser Val Val Tyr Leu Pro Glu Asn Leu Tyr Ser
50 55 60
Tyr Gly Trp Tyr Lys Gly Lys Thr Val Glu Pro Asn Gln Leu Ile Ala
65 70 75 80
Ala Tyr Val Ile Asp Asp Thr His Val Arg Thr Pro Gly Pro Ala Tyr
85 90 95
Ser Gly Arg Glu Thr Ile Ser Pro Ser Gly Asp Leu His Phe Gln Asn
100 105 110
Val Thr Leu Glu Asp Thr Gly Tyr Tyr Asn Leu Gln Val Thr Tyr Arg
115 120 125
Asn Ser Gln Ile Glu Gln Ala Ser His His Leu Arg Val Tyr Gln Val
130 135 140
Ser Gly Leu Thr Pro Pro Ser Lys Pro Ala Ala Pro Gln Ser Pro Arg
145 150 155 160
Arg Ala Pro Gly Val Leu Thr Cys His Thr Asn Asn Thr Gly Thr Ser
165 170 175
Phe Gln Trp Ile Phe Asn Asn Gln Arg Leu Gln Val Thr Lys Arg Met

```
            180                 185                 190
Lys Leu Ser Trp Phe Asn His Met Leu Thr Ile Asp Pro Ile Arg Gln
        195                 200                 205
Glu Asp Ala Gly Glu Tyr Gln Cys Glu Val Ser Asn Pro Val Ser Ser
    210                 215                 220
Asn Arg Ser Asp Pro Leu Lys Leu Thr Val Lys Ser Asp Asp Asn Thr
225                 230                 235                 240
Leu Gly Ile Leu Ile Gly Val Leu Val Gly Ser Leu Leu Val Ala Ala
                245                 250                 255
Leu Val Cys Phe Leu Leu Leu Arg Lys Thr Gly
            260                 265     267
```

<210> 1283
<211> 262
<212> PRT
<213> Homo sapiens

<400> 1283

```
Met Leu Val Leu Leu Val Leu Arg Val Ser Leu Ala Ala Leu Val Lys
 1               5                  10                  15
Met Glu Leu Leu Val Arg Trp Ala Pro Val Ala Cys Leu Val Arg Glu
            20                  25                  30
Val Ala Leu Glu Pro Leu Ala Leu Leu Val Leu Val Glu Met Met Val
        35                  40                  45
Leu Leu Val Leu Pro Gly Pro Leu Val Pro Pro Ala Pro Leu Val Leu
    50                  55                  60
Leu Ala Ser Leu Val Leu Leu Val Leu Arg Val Lys Leu Val Pro Lys
65                  70                  75                  80
Gly Pro Glu Ala Leu Lys Val Pro Arg Val Cys Val Val Ser Leu Ala
                85                  90                  95
Pro Leu Ala Leu Leu Val Leu Leu Ala Leu Leu Glu Thr Leu Val Leu
            100                 105                 110
Arg Glu Ser Leu Val Leu Lys Val Pro Met Val Leu Leu Val Leu Leu
        115                 120                 125
Val Leu Leu Ala Ser Leu Val Pro Glu Ala Pro Leu Asp Pro Arg Ala
    130                 135                 140
Pro Ala Ala Leu Leu Val Pro Arg Val Thr Ala Val Asn Leu Val Leu
145                 150                 155                 160
Leu Ala Ala Lys Glu Thr Leu Val Leu Arg Glu Ser Leu Ala Leu Leu
                165                 170                 175
Val Phe Lys Asp Pro Leu Ala Leu Leu Glu Arg Lys Glu Ser Glu Glu
            180                 185                 190
Leu Glu Val Asn Pro Asp Pro Leu Ala Cys Pro Asp Pro Leu Ala Ser
        195                 200                 205
Val Val Asp Leu Val Ala Val Val Ser Leu Ala Gln Met Val Leu Leu
    210                 215                 220
Val Pro Arg Val Pro Leu Val Asn Val Val Leu Leu Ala Leu Leu Ala
225                 230                 235                 240
Pro Lys Asp Leu Leu Val Lys Leu Val Val Pro Val Lys Leu Val Cys
                245                 250                 255
Leu Val Pro Arg Val  *
            260 261
```

<210> 1284

<211> 50
<212> PRT
<213> Homo sapiens

<400> 1284
Met Val Ile Leu Pro Leu Leu Leu Leu Ile Thr Thr Pro Pro Met Thr
1 5 10 15
Phe Leu Ala Phe Leu Leu Thr Leu Ile Leu Ser Cys Lys Asn Cys Ser
20 25 30
Lys Leu Ala Ala Ser Met Ile Arg Leu Leu Trp Gly Gly Cys Asn Gln
35 40 45
Glu *
49

<210> 1285
<211> 323
<212> PRT
<213> Homo sapiens

<400> 1285
Met Leu Val Met Ala Pro Arg Thr Val Leu Leu Leu Leu Ser Ala Ala
1 5 10 15
Leu Ala Leu Thr Glu Thr Trp Ala Gly Ser His Ser Met Arg Tyr Phe
20 25 30
Tyr Thr Ser Val Ser Arg Pro Gly Arg Gly Glu Pro Arg Phe Ile Ser
35 40 45
Val Gly Tyr Val Asp Asp Thr Gln Phe Val Arg Phe Asp Ser Asp Ala
50 55 60
Ala Ser Pro Arg Glu Glu Pro Arg Ala Pro Trp Ile Glu Gln Glu Gly
65 70 75 80
Pro Glu Tyr Trp Asp Arg Asn Thr Gln Ile Tyr Lys Ala Gln Ala Gln
85 90 95
Thr Asp Arg Glu Ser Leu Arg Asn Leu Arg Gly Tyr Tyr Asn Gln Ser
100 105 110
Glu Ala Gly Ser His Thr Leu Gln Ser Met Tyr Gly Cys Asp Val Gly
115 120 125
Pro Asp Gly Arg Leu Leu Arg Gly His Asp Gln Tyr Ala Tyr Asp Gly
130 135 140
Lys Asp Tyr Ile Ala Leu Asn Glu Asp Leu Arg Ser Trp Thr Ala Ala
145 150 155 160
Asp Thr Ala Ala Gln Ile Thr Gln Arg Lys Trp Glu Ala Ala Arg Glu
165 170 175
Ala Glu Gln Arg Arg Ala Tyr Leu Glu Gly Glu Cys Val Glu Trp Leu
180 185 190
Arg Arg Tyr Leu Glu Asn Gly Lys Asp Lys Leu Glu Arg Ala Asp Pro
195 200 205
Pro Lys Thr His Val Thr His His Pro Ile Ser Asp His Glu Ala Thr
210 215 220
Leu Arg Cys Trp Ala Leu Gly Phe Tyr Pro Ala Glu Ile Thr Leu Thr
225 230 235 240
Trp Gln Arg Asp Gly Glu Asp Gln Thr Gln Asp Thr Glu Leu Val Glu
245 250 255
Thr Arg Pro Ala Gly Asp Arg Thr Phe Gln Lys Val Gly Gln Leu Trp
260 265 270
Val Val Pro Ser Gly Glu Glu Gln Arg Tyr Thr Cys His Val Gln His

```
        275                 280                 285
Val Gly Ala Ala Glu Ala Pro His Pro Ser Glu Met Gly Ser Gly Leu
    290                 295                 300
Pro Ser Ser Thr Val Pro His Arg Trp Ala Leu Val Leu Gly Leu Gly
305                 310                 315                 320
Cys Pro  *
    322
```

```
     <210> 1286
     <211> 306
     <212> PRT
     <213> Homo sapiens

     <400> 1286
Met Leu Leu Phe Leu Leu Ser Ala Leu Val Leu Leu Thr Gln Pro Leu
 1               5                  10                  15
Gly Tyr Leu Glu Ala Glu Met Lys Thr Tyr Ser His Arg Thr Met Pro
            20                  25                  30
Ser Ala Cys Thr Leu Val Met Cys Ser Ser Val Glu Ser Gly Leu Pro
        35                  40                  45
Gly Arg Asp Gly Arg Asp Gly Arg Glu Gly Pro Arg Gly Glu Lys Gly
    50                  55                  60
Asp Pro Gly Leu Pro Gly Ala Ala Gly Gln Ala Gly Met Pro Gly Gln
65                  70                  75                  80
Ala Gly Pro Val Gly Pro Lys Gly Asp Asn Gly Ser Val Gly Glu Pro
                85                  90                  95
Gly Pro Lys Gly Asp Thr Gly Pro Ser Gly Pro Pro Gly Pro Pro Gly
            100                 105                 110
Val Pro Gly Pro Ala Gly Arg Glu Gly Pro Leu Gly Lys Gln Gly Asn
        115                 120                 125
Ile Gly Pro Gln Gly Lys Pro Gly Pro Lys Gly Glu Ala Gly Pro Lys
    130                 135                 140
Gly Glu Val Gly Ala Pro Gly Met Gln Gly Ser Ala Gly Ala Arg Gly
145                 150                 155                 160
Leu Ala Gly Pro Lys Gly Glu Arg Gly Val Pro Gly Glu Arg Gly Val
                165                 170                 175
Pro Gly Asn Thr Gly Ala Ala Gly Ser Ala Gly Ala Met Gly Pro Gln
            180                 185                 190
Gly Ser Pro Gly Ala Arg Gly Pro Pro Gly Leu Lys Gly Asp Lys Gly
        195                 200                 205
Ile Pro Gly Asp Lys Gly Ala Lys Gly Glu Ser Gly Leu Pro Asp Val
    210                 215                 220
Ala Ser Leu Arg Gln Gln Val Glu Ala Leu Gln Gly Gln Val Gln His
225                 230                 235                 240
Leu Gln Ala Ala Phe Ser Gln Tyr Lys Lys Val Glu Leu Phe Pro Asn
                245                 250                 255
Gly Gln Ser Val Gly Glu Lys Ile Phe Lys Thr Ala Gly Phe Val Lys
            260                 265                 270
Pro Phe Thr Glu Ala Gln Leu Leu Cys Thr Gln Ala Gly Gly Gln Leu
        275                 280                 285
Ala Ser Pro Arg Ser Ala Ala Glu Asn Ala Pro Leu Ala Thr Ala Gly
    290                 295                 300
Pro  *
305
```

<210> 1287
<211> 299
<212> PRT
<213> Homo sapiens

<400> 1287
Met Gly Arg Trp Ala Leu Asp Val Ala Phe Leu Trp Lys Ala Val Leu
1               5                   10                  15
Thr Leu Gly Leu Val Leu Leu Tyr Tyr Cys Phe Ser Ile Gly Ile Thr
            20                  25                  30
Phe Tyr Asn Lys Trp Leu Thr Lys Ser Phe His Phe Pro Leu Phe Met
        35                  40                  45
Thr Met Leu His Leu Ala Val Ile Phe Leu Phe Ser Ala Leu Ser Arg
    50                  55                  60
Ala Leu Val Gln Cys Ser Ser His Arg Ala Arg Val Val Leu Ser Trp
65                  70                  75                  80
Ala Asp Tyr Leu Arg Arg Val Ala Pro Thr Ala Leu Ala Thr Ala Leu
                85                  90                  95
Asp Val Gly Leu Ser Asn Trp Ser Phe Leu Tyr Val Thr Val Ser Leu
            100                 105                 110
Tyr Thr Met Thr Lys Ser Ser Ala Val Leu Phe Ile Leu Ile Phe Ser
        115                 120                 125
Leu Ile Phe Lys Leu Glu Glu Leu Arg Ala Ala Leu Val Leu Val Val
    130                 135                 140
Leu Leu Ile Ala Gly Gly Leu Phe Met Phe Thr Tyr Lys Ser Thr Gln
145                 150                 155                 160
Phe Asn Val Glu Gly Phe Ala Leu Val Leu Gly Ala Ser Phe Ile Gly
                165                 170                 175
Gly Ile Arg Trp Thr Leu Thr Gln Met Leu Leu Gln Lys Ala Glu Leu
            180                 185                 190
Gly Leu Gln Asn Pro Ile Asp Thr Met Phe His Leu Gln Pro Leu Met
        195                 200                 205
Phe Leu Gly Leu Phe Pro Leu Phe Ala Val Phe Glu Gly Leu His Leu
    210                 215                 220
Ser Thr Ser Glu Lys Ile Phe Arg Phe Gln Gly His Arg Ala Ala Pro
225                 230                 235                 240
Ala Gly Thr Trp Gly Ala Ser Ser Leu Ala Gly Phe Ser Pro Leu Val
                245                 250                 255
Trp Ala Ser Leu Ser Ser Ser Trp Ser Pro Glu Pro Pro Ala Ser Leu
            260                 265                 270
Ser Pro Leu Pro Ala Phe Leu Arg Lys Ser Ala Leu Cys Cys Trp Gln
        275                 280                 285
Leu Ile Cys Trp Ala Ile Arg Ser Ala Ser *
    290                 295         298

<210> 1288
<211> 161
<212> PRT
<213> Homo sapiens

<400> 1288
Met Glu Ser Ala Leu Pro Ala Ala Gly Phe Leu Tyr Trp Val Gly Ala
1               5                   10                  15
Gly Thr Val Ala Tyr Leu Ala Leu Arg Ile Ser Tyr Ser Leu Phe Thr

20 25 30
Ala Leu Arg Val Trp Gly Val Gly Asn Glu Ala Gly Val Gly Pro Gly
35 40 45
Leu Gly Glu Trp Ala Val Val Thr Gly Ser Thr Asp Gly Ile Gly Lys
50 55 60
Ser Tyr Ala Glu Glu Leu Ala Lys His Gly Met Lys Val Val Leu Ile
65 70 75 80
Ser Arg Ser Lys Asp Lys Leu Asp Gln Val Ser Ser Glu Ile Lys Glu
85 90 95
Lys Phe Lys Val Glu Thr Arg Thr Ile Ala Val Asp Phe Ala Ser Glu
100 105 110
Asp Ile Tyr Asp Lys Ile Lys Thr Gly Leu Ala Gly Leu Glu Ile Gly
115 120 125
Ile Leu Val Asn Asn Val Gly Met Ser Tyr Glu Tyr Pro Glu Tyr Phe
130 135 140
Leu Asp Val Pro Asp Leu Asp Asn Val Ile Lys Lys Asn Asp Lys Tyr
145 150 155 160
*

<210> 1289
<211> 46
<212> PRT
<213> Homo sapiens

<400> 1289
Met Val Leu Ser Ala Pro Ser Leu Trp Pro Cys Ser Ser Phe Ser Ile
1 5 10 15
Ser Cys Leu His Val Gly Leu Thr Ala Phe Leu Phe Gln Val Ala Phe
20 25 30
Leu Cys Leu Leu Cys Cys Val Glu Leu Leu Leu Asp Val *
35 40 45

<210> 1290
<211> 453
<212> PRT
<213> Homo sapiens

<400> 1290
Met Thr Ser Lys Phe Ile Leu Val Ser Phe Ile Leu Ala Ala Leu Ser
1 5 10 15
Leu Ser Thr Thr Phe Ser Leu Gln Pro Asp Gln Gln Lys Val Leu Leu
20 25 30
Val Ser Phe Asp Gly Phe Arg Trp Asp Tyr Leu Tyr Lys Val Pro Thr
35 40 45
Pro His Phe His Tyr Ile Met Lys Tyr Gly Val His Val Lys Gln Val
50 55 60
Thr Asn Val Phe Ile Thr Lys Thr Tyr Pro Asn His Tyr Thr Leu Val
65 70 75 80
Thr Gly Leu Phe Ala Glu Asn His Gly Ile Val Ala Asn Asp Met Phe
85 90 95
Asp Pro Ile Arg Asn Lys Ser Phe Ser Leu Asp His Met Asn Ile Tyr
100 105 110

```
Asp Ser Lys Phe Trp Glu Glu Ala Thr Pro Ile Trp Ile Thr Asn Gln
        115                 120                 125
Arg Ala Gly His Thr Ser Gly Ala Ala Met Trp Pro Gly Thr Asp Val
    130                 135                 140
Lys Ile His Lys Arg Phe Pro Thr His Tyr Met Pro Tyr Asn Glu Ser
145                 150                 155                 160
Val Ser Phe Glu Asp Arg Val Ala Lys Ile Ile Glu Trp Phe Thr Ser
                165                 170                 175
Lys Glu Pro Ile Asn Leu Gly Leu Leu Tyr Trp Glu Asp Pro Asp Asp
            180                 185                 190
Met Gly His His Leu Gly Pro Asp Ser Pro Leu Met Gly Pro Val Ile
        195                 200                 205
Ser Asp Ile Asp Lys Lys Leu Gly Tyr Leu Ile Gln Met Leu Lys Lys
    210                 215                 220
Ala Lys Leu Trp Asn Thr Leu Asn Leu Ile Ile Thr Ser Asp His Gly
225                 230                 235                 240
Met Thr Gln Cys Ser Glu Glu Arg Leu Ile Glu Leu Asp Gln Tyr Leu
                245                 250                 255
Asp Lys Asp His Tyr Thr Leu Ile Asp Gln Ser Pro Val Ala Ala Ile
            260                 265                 270
Leu Pro Lys Glu Gly Lys Phe Asp Glu Val Tyr Glu Ala Leu Thr His
        275                 280                 285
Ala His Pro Asn Leu Thr Val Tyr Lys Lys Glu Asp Val Pro Glu Arg
    290                 295                 300
Trp His Tyr Lys Tyr Asn Ser Arg Ile Gln Pro Ile Ile Ala Val Ala
305                 310                 315                 320
Asp Glu Gly Trp His Ile Leu Gln Asn Lys Ser Asp Asp Phe Leu Leu
                325                 330                 335
Gly Asn His Gly Tyr His Asn Ala Leu Ala Asp Met His Pro Ile Phe
            340                 345                 350
Leu Ala His Gly Pro Ala Phe Arg Lys Asn Phe Ser Lys Glu Ala Met
        355                 360                 365
Asn Ser Thr Asp Leu Tyr Pro Leu Leu Cys His Leu Leu Asn Ile Thr
    370                 375                 380
Ala Met Pro His Asn Gly Ser Phe Trp Asn Val Gln Asp Leu Leu Asn
385                 390                 395                 400
Ser Ala Met Pro Arg Val Val Pro Tyr Thr Gln Ser Thr Ile Leu Leu
                405                 410                 415
Pro Gly Ser Val Lys Pro Ala Glu Tyr Asp Gln Glu Gly Ser Tyr Pro
            420                 425                 430
Tyr Phe Ile Gly Val Ser Leu Gly Ser Ile Ile Val Ile Val Phe Phe
        435                 440                 445
Cys Asn Phe His  *
    450     452
```

<210> 1291
<211> 78
<212> PRT
<213> Homo sapiens

<221> misc_feature
<222> (1)...(78)
<223> Xaa = any amino acid or nothing

<400> 1291

```
Met Leu Ser Val Thr Ala Phe Ile Leu Ala Glu Thr Val Leu Ala Ser
```

```
 1               5                  10                  15
Gln Glu Val Gln Gly Gly Val Gln Val Arg Val Tyr Leu Met Asn Ala
            20                  25                  30
Val Pro Asp Gly Leu Gln Gly Gly Ser Pro Val Gly Gly Leu Gly Leu
        35                  40                  45
Leu Leu Ala Pro Asp Asn Ser Gly His Arg Arg Ser Ser Cys Arg Ile
    50                  55                  60
Pro Ala Ala Arg Val Tyr Xaa Xaa Xaa Xaa Pro Arg Pro Pro
65                  70                  75          78


     <210> 1292
     <211> 416
     <212> PRT
     <213> Homo sapiens


     <400> 1292
Met Val Leu Trp Ile Leu Trp Arg Pro Phe Gly Phe Ser Gly Arg Phe
 1               5                  10                  15
Leu Lys Leu Glu Ser His Ser Ile Thr Glu Ser Lys Ser Leu Ile Pro
            20                  25                  30
Val Ala Trp Thr Ser Leu Thr Gln Met Leu Leu Glu Ala Pro Gly Ile
        35                  40                  45
Phe Leu Leu Gly Gln Arg Lys Arg Phe Ser Thr Met Pro Glu Thr Glu
    50                  55                  60
Thr His Glu Arg Glu Thr Glu Leu Phe Ser Pro Pro Ser Asp Val Arg
65                  70                  75                  80
Gly Met Thr Lys Leu Asp Arg Thr Ala Phe Lys Lys Thr Val Asn Ile
                85                  90                  95
Pro Val Leu Lys Val Arg Lys Glu Ile Val Ser Lys Leu Met Arg Ser
            100                 105                 110
Leu Lys Arg Ala Ala Leu Gln Arg Pro Gly Ile Arg Arg Val Ile Glu
        115                 120                 125
Asp Pro Glu Asp Lys Glu Ser Arg Leu Ile Met Leu Asp Pro Tyr Lys
    130                 135                 140
Ile Phe Thr His Asp Ser Phe Glu Lys Ala Glu Leu Ser Val Leu Glu
145                 150                 155                 160
Gln Leu Asn Val Ser Pro Gln Ile Ser Lys Tyr Asn Leu Glu Leu Thr
                165                 170                 175
Tyr Glu His Phe Lys Ser Glu Glu Ile Leu Arg Ala Val Leu Pro Glu
            180                 185                 190
Gly Gln Asp Val Thr Ser Gly Phe Ser Arg Ile Gly His Ile Ala His
        195                 200                 205
Leu Asn Leu Arg Asp His Gln Leu Pro Phe Lys His Leu Ile Gly Gln
    210                 215                 220
Val Met Ile Asp Lys Asn Pro Gly Ile Thr Ser Ala Val Asn Lys Ile
225                 230                 235                 240
Asn Asn Ile Asp Asn Met Tyr Arg Asn Phe Gln Met Glu Val Leu Ser
                245                 250                 255
Gly Glu Gln Asn Met Met Thr Lys Val Arg Glu Asn Asn Tyr Thr Tyr
            260                 265                 270
Glu Phe Asp Phe Ser Lys Val Tyr Trp Asn Pro Arg Leu Ser Thr Glu
        275                 280                 285
His Ser Arg Ile Thr Glu Leu Leu Lys Pro Gly Asp Val Leu Phe Asp
    290                 295                 300
Val Phe Ala Gly Val Gly Pro Phe Ala Ile Pro Val Ala Lys Lys Asn
305                 310                 315                 320
```

Cys Thr Val Phe Ala Asn Asp Leu Asn Pro Glu Ser His Lys Trp Leu
325 330 335
Leu Tyr Asn Cys Lys Leu Asn Lys Val Asp Gln Lys Val Lys Val Phe
340 345 350
Asn Leu Asp Gly Lys Asp Phe Leu Gln Gly Pro Val Lys Glu Glu Leu
355 360 365
Met Gln Leu Leu Gly Leu Ser Lys Glu Arg Lys Pro Ser Val His Val
370 375 380
Val Met Asn Leu Pro Ala Lys Ala Ile Glu Phe Leu Ser Ala Phe Lys
385 390 395 400
Trp Leu Leu Asp Gly Gln Pro Met Pro Ala Val Ser Ser Phe Pro *
405 410 415

<210> 1293
<211> 113
<212> PRT
<213> Homo sapiens

<400> 1293
Met Val Arg Pro Leu Leu Leu Leu Asn Leu His Phe His Leu Pro Ser
1 5 10 15
Leu Val Ser Leu Ser Leu Ser Leu Leu Leu Ser Val Ser Leu Ser Leu
20 25 30
Val Asn Ala Val Arg Leu Leu Arg Ala Ser Phe Cys Ser Trp Leu Ile
35 40 45
Ala Lys Ser Leu Ile Thr Leu Trp Val Arg Pro Ser Gln Ile Gly Lys
50 55 60
Leu Lys Ala Leu Ala Ser Ser Thr Thr Ser Met Ala Trp Glu Gly Leu
65 70 75 80
Leu Asp Thr Phe Ala Leu Ser Ile Ser Ser Phe Ser Asn Ser Leu Leu
85 90 95
Gly Ile Leu Leu Cys Phe Leu Lys Ser Pro Asn Ile Phe Gln Ala Ser
100 105 110 112
*

<210> 1294
<211> 57
<212> PRT
<213> Homo sapiens

<400> 1294
Met Asp Phe Leu Met Leu Ala Val Cys Ala His Arg Leu Cys Phe Leu
1 5 10 15
Tyr Leu Phe Ile Leu Tyr Glu Ser Lys Asn Lys Arg Glu Cys Glu Gln
20 25 30
Phe Arg Arg Leu Gln Ile Tyr Leu Val Arg Leu Leu Ser Lys Arg Phe
35 40 45
Pro Val Val Val Ile Pro Ala Val *
50 55 56

<210> 1295
<211> 68
<212> PRT
<213> Homo sapiens

<400> 1295
Met Phe Leu Ser Leu Cys Leu Leu Ser Ala Ala Leu Thr Lys Ile Ser
1 5 10 15
Ser Lys Ile Leu Tyr Lys Pro Gly Thr Lys Val Thr Ser Leu Gln Phe
20 25 30
Ile Pro Thr Ser Ser Ser Tyr Thr His Met Asn Cys Val Asn Gly Ser
35 40 45
Thr Asp Pro Ile Tyr Val Ser Gly Arg Arg Arg Met Cys Ser Ser Cys
50 55 60
Val Phe Ile *
65 67

<210> 1296
<211> 66
<212> PRT
<213> Homo sapiens

<400> 1296
Met Trp Ser Ala His Pro Leu Ala Val Leu Ser Leu Lys Leu Thr Leu
1 5 10 15
Phe Ser Leu Thr Ser Asp Trp Leu Ser Ser Lys Asp Met Ala Ile Ser
20 25 30
Leu Ala Phe Lys Ile Ser Gln Ile Leu Cys Ser Val Leu Ser Ala Pro
35 40 45
Gly Lys Arg Leu Ile Ser Val Leu Trp Asn Thr Ser Ser Leu Lys Arg
50 55 60
Ser *
65

<210> 1297
<211> 57
<212> PRT
<213> Homo sapiens

<400> 1297
Met Leu His Ser Gln Leu Leu Ala Val Ser Phe Arg Leu Ile Val Thr
1 5 10 15
Leu Pro Leu Ser Ile Gln Asp Trp Asp Asp Ala Glu Asn Met Lys Gly
20 25 30
Leu Gln Tyr Ile Phe Asn Thr Leu Trp Ser Val Ser Ser Pro Val Ile
35 40 45
Thr Ser Ile Leu Ser Ser Lys His *
50 55 56

<210> 1298

<211> 235
<212> PRT
<213> Homo sapiens

<400> 1298
Met Arg Lys Thr Arg Leu Trp Gly Leu Leu Trp Met Leu Phe Val Ser
1 5 10 15
Glu Leu Arg Ala Ala Thr Lys Leu Thr Glu Glu Lys Tyr Glu Leu Lys
20 25 30
Glu Gly Gln Thr Leu Asp Val Lys Cys Asp Tyr Thr Leu Glu Lys Phe
35 40 45
Ala Ser Ser Gln Lys Ala Trp Gln Ile Ile Arg Asp Gly Glu Met Pro
50 55 60
Lys Thr Leu Ala Cys Thr Glu Arg Pro Ser Lys Asn Ser His Pro Val
65 70 75 80
Gln Val Gly Arg Ile Ile Leu Glu Asp Tyr His Asp His Gly Leu Leu
85 90 95
Arg Val Arg Met Val Asn Leu Gln Val Glu Asp Ser Gly Leu Tyr Gln
100 105 110
Cys Val Ile Tyr Gln Pro Pro Lys Glu Pro His Met Leu Phe Asp Arg
115 120 125
Ile Arg Leu Val Val Thr Lys Gly Phe Ser Gly Thr Pro Gly Ser Asn
130 135 140
Glu Asn Ser Thr Gln Asn Val Tyr Lys Ile Pro Pro Thr Thr Thr Lys
145 150 155 160
Ala Leu Cys Pro Leu Tyr Thr Thr Pro Arg Thr Val Thr Gln Ala Pro
165 170 175
Pro Lys Ser Thr Ala Asp Val Ser Thr Pro Asp Ser Glu Ile Asn Leu
180 185 190
Thr Asn Val Thr Asp Ile Ile Arg Val Pro Val Phe Asn Ile Val Ile
195 200 205
Leu Leu Ala Gly Gly Phe Leu Ser Lys Ser Leu Val Phe Ser Val Leu
210 215 220
Phe Ala Val Thr Leu Arg Ser Phe Val Pro *
225 230 234

<210> 1299
<211> 64
<212> PRT
<213> Homo sapiens

<400> 1299
Met Arg Trp Lys Val Gln Val Asn Ser Leu Met Val Leu Pro Ser Leu
1 5 10 15
Thr Val Cys Tyr Ser Thr His Leu Ser Thr Gly Cys Arg His Ile Lys
20 25 30
Val Asn Val Gln Val Leu Glu Asn Ile Gln Arg Ile Leu Asn Val Gln
35 40 45
Asn Ser Glu Lys Gln Ile Tyr Ala Glu Cys Val Val Gly Ala Phe *
50 55 60 63

<210> 1300
<211> 80

<212> PRT
<213> Homo sapiens

<400> 1300
Met Ala Ser Arg Ser Asn Tyr Leu Thr Glu Thr Leu Thr Pro Phe Pro
1 5 10 15
Ala Leu Leu Ser Leu Phe Met Leu Tyr Leu Ser His Thr Gly Phe Asp
20 25 30
Asn Ile Ile Pro Thr Phe Pro Thr Lys Pro Ala Tyr Thr Leu His Arg
35 40 45
Leu Leu Pro His Cys Pro Asp Ile His Ile Ala Tyr Ser Leu Ile Ser
50 55 60
Ser His Leu Phe Ala Gln Gly Ala Ser Leu Ser Thr Arg Thr His *
65 70 75 79

<210> 1301
<211> 87
<212> PRT
<213> Homo sapiens

<400> 1301
Met Arg Phe Arg Ala Glu Pro Lys Ser Arg Pro Leu Pro Ala Leu Cys
1 5 10 15
His Val Leu Ile Ala Cys Ile Val Phe Arg Trp Ala Phe Ala Gln Pro
20 25 30
Leu Pro Ser Ser Arg Ser Tyr Arg Ser Ser Gly Glu Phe Pro Arg Ser
35 40 45
Pro Ser Phe Lys Lys Thr Lys Thr Pro Ser Trp Gly Glu Arg Arg Val
50 55 60
Leu Leu Tyr Ser Arg Met Leu Arg Ala Asn Leu Arg Met Trp Arg Glu
65 70 75 80
Tyr Trp Ser Gln Lys Ser Ile
85 87

<210> 1302
<211> 143
<212> PRT
<213> Homo sapiens

<400> 1302
Met Asp His Cys Gly Ala Leu Phe Leu Cys Leu Cys Leu Leu Thr Leu
1 5 10 15
Gln Asn Ala Thr Thr Glu Thr Trp Glu Glu Leu Leu Ser Tyr Met Glu
20 25 30
Asn Met Gln Val Ser Arg Gly Arg Ser Ser Val Phe Ser Ser Arg Gln
35 40 45
Leu His Gln Leu Glu Gln Met Leu Leu Asn Thr Ser Phe Pro Gly Tyr
50 55 60
Asn Leu Thr Leu Gln Thr Pro Thr Ile Gln Ser Leu Ala Phe Lys Leu
65 70 75 80
Ser Cys Asp Phe Ser Gly Leu Ser Leu Thr Ser Ala Thr Leu Lys Arg
85 90 95

Val Pro Gln Ala Gly Gly Gln His Ala Arg Gly Gln His Ala Met Gln
100 105 110
Phe Pro Ala Glu Leu Thr Arg Asp Ala Cys Lys Thr Arg Pro Arg Glu
115 120 125
Leu Arg Leu Ile Cys Ile Tyr Phe Ser Asn Thr His Phe Phe Lys
130 135 140 143

<210> 1303
<211> 60
<212> PRT
<213> Homo sapiens

<400> 1303
Met Ile Leu Leu Met Ser Ala Ala Ile Phe Cys Ser Ala Glu Val Phe
1 5 10 15
Thr Arg Gly Ser Phe Phe Ser Asp Met Leu Thr Leu Asp Arg Val Lys
20 25 30
Ala Lys Gly Leu Gln Gly Glu Gly Ala Ala Ser Thr Cys Ala Leu Ala
35 40 45
Ala Asp Ser Gln Gly Ser Gly Ala Ser Gly Thr Lys
50 55 60

<210> 1304
<211> 56
<212> PRT
<213> Homo sapiens

<400> 1304
Met Lys Met Met Phe Ile Ile Thr Asn Trp Leu Asn Tyr Tyr Phe Leu
1 5 10 15
Leu Phe Ser Pro Ser Asn Pro Gln Ile Gln Ser Ile Leu His Glu Val
20 25 30
Ala Pro Leu Trp Phe Arg Thr Leu Tyr Thr Leu Leu Arg Gly Cys Ser
35 40 45
Thr Trp Lys Gly Leu Ser Ser *
50 55

<210> 1305
<211> 63
<212> PRT
<213> Homo sapiens

<400> 1305
Met Asn Ile Ile Phe Ile Tyr Leu Ala Thr Ser Leu Ala Phe Leu Ile
1 5 10 15
Ile Asn Leu Ser Gln Leu Leu Phe Thr Glu Tyr Leu His Phe Arg Cys
20 25 30
Cys Ser Lys Cys Ser Thr Cys Ile Asn Leu Leu Ser His His Glu Trp
35 40 45
Glu Leu Leu Pro Ser Ser Tyr Arg Arg Gly Ser Arg Ser Pro *

50 55 60 62

<210> 1306
<211> 138
<212> PRT
<213> Homo sapiens

<400> 1306
```
Met Gln Asn Arg Thr Gly Leu Ile Leu Cys Ala Leu Ala Leu Leu Met
 1               5                  10                  15
Gly Phe Leu Met Val Cys Leu Gly Ala Phe Phe Ile Ser Trp Gly Ser
            20                  25                  30
Ile Phe Asp Cys Gln Gly Ser Leu Ile Ala Ala Tyr Leu Leu Leu Pro
        35                  40                  45
Leu Gly Phe Val Ile Leu Leu Ser Gly Ile Phe Trp Ser Asn Tyr Arg
    50                  55                  60
Gln Val Thr Glu Ser Lys Gly Val Leu Arg His Met Leu Arg Gln His
65                  70                  75                  80
Leu Ala His Gly Ala Leu Pro Val Ala Thr Val Asp Arg Pro Asp Phe
                85                  90                  95
Tyr Pro Pro Ala Tyr Glu Glu Ser Leu Glu Val Glu Lys Gln Ser Cys
            100                 105                 110
Pro Ala Glu Arg Glu Ala Pro Arg His Ser Ser Thr Ser Ile Tyr Arg
        115                 120                 125
Asp Gly Pro Gly Ile Pro Gly Trp Lys  *
    130                 135     137
```

<210> 1307
<211> 64
<212> PRT
<213> Homo sapiens

<400> 1307
```
Met Met Ala Ile Lys Pro Thr Ile Leu Val Thr Gln Gly Leu Ile Leu
 1               5                  10                  15
Cys Trp Lys Cys His Lys Met Ile Cys Ser Tyr Phe Asn Leu Gln Leu
            20                  25                  30
Glu Arg His Phe Leu Glu Thr Ile Gln Ser Asp Ser Phe Met Glu Lys
        35                  40                  45
Leu Thr Leu Thr Asp Leu Thr Ile Tyr Arg Ile His Val Ala Thr His
    50                  55                  60                  64
```

<210> 1308
<211> 65
<212> PRT
<213> Homo sapiens

<400> 1308

```
Met Pro Cys Ser Gly Ser Ser Val Gln Thr Phe Arg Pro Leu Leu Ile
 1               5                  10                  15
Phe His Asn Val Thr Phe Phe Ile Leu Pro Val Lys Cys Phe Asn Ala
            20                  25                  30
Leu Ile Asn Val Leu Glu Arg Pro Phe Trp Gln Leu Leu Gly Glu Ile
        35                  40                  45
Gly Glu Glu Tyr Arg Gly Ser Glu Asp Trp Leu Gly Gly Ser Phe Arg
    50                  55                  60              64
 *
```

<210> 1309
<211> 75
<212> PRT
<213> Homo sapiens

<400> 1309

```
Met Arg Ile Trp His Arg Trp Leu Leu Val Arg Ile Leu Phe Pro Ala
 1               5                  10                  15
Pro Gly Leu Gln Thr Ala Thr Phe Ser Val Cys Phe His Val Ala Glu
            20                  25                  30
Ser Glu Leu Trp His Leu Leu Cys Phe Phe Phe Phe Leu Ala Leu Leu
        35                  40                  45
Pro Pro Arg Trp Lys Ala Arg Gly Pro Ile Trp Val His Gly Thr Leu
    50                  55                  60
Gly Phe Arg Val Gly Arg Asn Phe Leu Ala  *
65                  70              74
```

<210> 1310
<211> 46
<212> PRT
<213> Homo sapiens

<400> 1310

```
Met Lys Leu Gly Asp Val Phe Val Lys Leu Leu Val Ser Leu Ala Gly
 1               5                  10                  15
Glu Ile Leu Leu Ala Pro Leu Val Ser Ala Ser Gly Met Gly Pro Ala
            20                  25                  30
Gly Val Glu Ala Leu Glu Glu Val Ser Ala Leu Ser Val  *
        35                  40                  45
```

<210> 1311
<211> 105
<212> PRT
<213> Homo sapiens

<400> 1311

```
Met Tyr Trp Val Thr Val Ile Thr Leu Ile Tyr Gly Tyr Tyr Ala Trp
 1               5                  10                  15
Val Gly Phe Trp Pro Glu Ser Ile Pro Tyr Gln Asn Leu Gly Pro Leu
```

```
                20                  25                  30
Gly Pro Leu Thr Gln Tyr Leu Met Asp His His His Thr Leu Leu Cys
            35                  40                  45
Asn Gly Tyr Trp Leu Ala Trp Leu Ile His Val Gly Glu Ser Leu His
        50                  55                  60
Ala Ile Leu Leu Gly Glu Arg Lys Gly Ile Thr Ser Gly Arg Ser Gln
65                  70                  75                  80
Leu Leu Trp Leu Leu Gln Thr Leu Phe Phe Gly Ile Thr Thr Leu Thr
                85                  90                  95
Ile Phe Asp Ala Tyr Lys Arg Lys Arg
            100                 105
```

```
<210> 1312
<211> 114
<212> PRT
<213> Homo sapiens

<400> 1312
Met Lys Gly Lys Trp Cys Cys Ser Leu Leu Cys Gln Ser Pro Gln Val
 1               5                  10                  15
Gln Thr Ala Leu Val Cys Pro Leu Ser Leu Ser Leu Gly Pro Pro Gly
            20                  25                  30
Pro Gln Cys Pro Leu Leu Trp Leu Gly Gln Glu Asp Leu Pro Asp Ile
        35                  40                  45
Ala Arg Cys Ile Thr Asp Asp Cys Ser Gln Leu Pro Gln Ala Pro Ala
    50                  55                  60
Ser Leu Ala Ser Cys Phe Phe Pro Gln Ser Cys Leu Leu Ile Ser Ile
65                  70                  75                  80
His Leu Ser Met Gly Tyr Ser Trp Thr Leu Gly Leu Gly Val Gly Ile
                85                  90                  95
Arg Leu Leu Pro Thr Lys Gly Val Lys Val Thr His Phe Pro Tyr His
            100                 105                 110
Ala  *
113
```

```
<210> 1313
<211> 88
<212> PRT
<213> Homo sapiens

<400> 1313
Met Ser Ser Ser Gly Gln Leu Gly His Pro Pro Arg Ala Pro His Ser
 1               5                  10                  15
Trp Arg Arg Trp Cys Trp Trp Leu Phe Met Leu Ala Thr Ser Leu Ser
            20                  25                  30
Arg Arg Arg Arg Pro Ser Thr Pro Leu Ile His Tyr Arg Val Phe Thr
        35                  40                  45
Val Asn His Lys Met Asp Pro Val Thr Arg Thr Phe Thr Leu Asp Ile
    50                  55                  60
Lys Val Val Phe Pro Asp Glu Gly Trp Gly Val Val Val Asp Pro Gly
65                  70                  75                  80
His Trp Gly Tyr Met Val Cys  *
                85          87
```

<210> 1314
<211> 65
<212> PRT
<213> Homo sapiens

<400> 1314
Met Gly Gly Arg Leu Trp Ile Phe Leu Gln Leu Cys Gln Ser Leu Gly
1 5 10 15
Leu Ser Thr Val Val Ser Ser Arg Pro Val Ala Cys Leu Glu Ser Val
20 25 30
Pro Gly Met Cys Met Ser Val Cys Met Pro Leu Asn Tyr Arg Gly Ser
35 40 45
Asn Phe Ser Glu Thr Asp Val Trp Met Asp Leu Ser Arg Ala His Leu
50 55 60 64
*

<210> 1315
<211> 71
<212> PRT
<213> Homo sapiens

<400> 1315
Met Leu Ile Pro Ile Pro Val His Ile Phe Pro Leu Ser Ser Leu Leu
1 5 10 15
Gly Asp Gly Thr Met Arg Leu Leu Pro Asp Ile Ser Ser Asp Trp Leu
20 25 30
Cys Leu Asn Gln Glu Phe Ala Pro Val Gln Ser Ala Ile Ala Met Glu
35 40 45
Trp Gly Ser Cys Val Gly Asp Gln Asp Asp Thr His Trp Ile Cys Leu
50 55 60
Arg Gln Thr Ser Gly Val *
65 70

<210> 1316
<211> 114
<212> PRT
<213> Homo sapiens

<400> 1316
Met Ala Thr Pro Ser Ser Pro Trp Trp Ala His Ser Gly Leu Pro Pro
1 5 10 15
Leu Phe Ser Ser Gly Leu Ser Trp Arg Leu Val Pro Leu Phe Trp Cys
20 25 30
Leu Gln Ser Leu Thr Gly Phe Leu Gly Pro Cys Leu Pro Arg Thr Thr
35 40 45
Arg Ala Phe Leu Ser Leu Gln Ser Trp Asp Leu Pro Gly Thr Arg Pro
50 55 60
Gly Ser Gln Ala Gln Gly Phe Thr Ala Cys Asn Ala Ala Asn Thr Pro

```
 65                  70                  75                  80
Gly Leu Ala Ala Leu Pro Gly Ser Gly Ala Phe Ser Val Ile Pro Val
                85                  90                  95
Ser Leu Leu Leu Pro Val Pro Glu Gly Leu Gly Arg Thr Tyr Leu Tyr
            100                 105                 110
Ser  *
113


     <210> 1317
     <211> 91
     <212> PRT
     <213> Homo sapiens


     <400> 1317
Met Met Val Trp Asn Leu Phe Pro Cys Phe Pro Pro Leu Leu Leu Leu
  1               5                  10                  15
Gln Phe Ile Asp Cys Gln Gln Ser Ser Glu Ile Glu Gln Gly Phe Thr
            20                  25                  30
Arg Ser Leu Leu Gly His Pro Ile Phe Phe Cys Pro Asp Pro Cys Trp
        35                  40                  45
Gln Ser Cys Met Asn Cys Val Ile Leu Leu Ser Ala Phe Phe Phe Leu
    50                  55                  60
Phe Asp Lys Met Asp Ile Lys Asn Ser Cys Cys Ala Lys Val Ser Ser
 65                  70                  75                  80
Leu Leu Gln Glu Glu Asn Gln Phe Phe Phe  *
                85                  90


     <210> 1318
     <211> 65
     <212> PRT
     <213> Homo sapiens


     <400> 1318
Met Leu Pro Leu Ile Ser Ser Ile Lys Ile Leu Lys Leu Leu Tyr Tyr
  1               5                  10                  15
Phe Ser Val Trp Gly Trp Gly Phe Phe Phe Phe Glu Thr Glu Phe Arg
            20                  25                  30
Ser Cys Cys Pro Gly Trp Ser Ala Met Val Arg Ser Gln Leu Thr Ala
        35                  40                  45
Thr Ser Thr Ser Arg Val Gln Ala Ile Leu Leu Pro Gln Pro Pro Glu
    50                  55                  60              64
 *


     <210> 1319
     <211> 46
     <212> PRT
     <213> Homo sapiens


     <400> 1319
```

Met Val Thr Leu Leu Ile Ala Lys Gln Phe Trp Ile Phe Thr Val Asp
1 5 10 15
Leu His Leu Ser Asp Tyr Val Leu Glu Leu Ser Arg Tyr Leu Ile Asn
20 25 30
Ala Cys Phe Tyr Ser Pro Cys Ser Gln Pro Ile Glu Lys *
35 40 45

<210> 1320
<211> 47
<212> PRT
<213> Homo sapiens

<400> 1320
Met Pro Ala Leu Leu Val Leu Lys Val Val Lys Val Leu Leu Pro Met
1 5 10 15
Val Leu Thr Gly Leu Gly Val Glu Glu Leu Lys Glu Met Val Leu Leu
20 25 30
Leu Pro Val Pro Cys Ala Ala Ile Ile Gly Ser Phe Lys Leu *
35 40 45 46

<210> 1321
<211> 55
<212> PRT
<213> Homo sapiens

<400> 1321
Met Ile Cys Phe Cys Leu Pro Val Cys Pro Lys Thr His Leu Ala His
1 5 10 15
Pro Met Leu Ala Thr Leu Ala Phe Val Ser Leu Leu Glu Tyr Ala Lys
20 25 30
His Cys Leu Arg Asp Phe Ile Leu Val Ser Phe Leu Leu Gly Met Leu
35 40 45
Phe Leu Arg Tyr Gln His *
50 54

<210> 1322
<211> 301
<212> PRT
<213> Homo sapiens

<400> 1322
Met Lys Ile Ala Phe Gly Asn Leu Trp Met Glu Ile Leu Tyr Leu Lys
1 5 10 15
Pro Pro Trp Thr Leu Leu His Leu Leu Gln Cys Phe Lys Lys His Trp
20 25 30
Leu Ala Val Phe Gly Leu Val Met Glu Lys Asn Leu Leu Leu Thr Ile
35 40 45
Glu Ser Leu Tyr Lys Asn Leu Arg Lys Ala Asn Lys Ala Val Asp Phe
50 55 60
Thr Thr Val Lys Phe Leu Leu Gln Asp Ser Arg Ser Leu Leu His Ala

```
 65                  70                  75                  80
Phe Ser Thr Arg Ser Asn Tyr Asp Gly Ile Leu Pro Gln Thr Phe Ala
                 85                  90                  95
Gln Val Asn Asn Leu Leu Gln Thr Phe Ala Glu Val Lys Thr Lys Leu
            100                 105                 110
Lys Pro Asn Ser Ser Glu Asn Thr Val Thr Lys Lys Gln Glu Gly Thr
        115                 120                 125
Ser Leu Lys Asn Ser His Asn Gln Glu Ile Thr Val Phe Ser Ser Ser
    130                 135                 140
His Leu Pro Gln Pro Ser Arg His Gln Glu Ile Trp Ser Ile Leu Glu
145                 150                 155                 160
Ser Val Trp Ile Thr Ile Tyr Gln Asn Ser Thr Asp Val Phe Gln Arg
                165                 170                 175
Leu Gly Ser Asn Ser Ala Leu Thr Thr Ser Asn Ile Ala Ser Phe Glu
            180                 185                 190
Glu Ala Phe Ile Cys Leu Gln Lys Leu Met Ala Ala Val Arg Asp Ile
        195                 200                 205
Leu Glu Gly Ile Gln Arg Ile Leu Ala Pro Asn Ser Asn Tyr Gln Asp
    210                 215                 220
Val Glu Thr Leu Tyr Asn Phe Leu Ile Lys Tyr Glu Val Asn Lys Asn
225                 230                 235                 240
Val Lys Phe Thr Ala Gln Glu Ile Tyr Asp Cys Val Ser Gln Thr Glu
                245                 250                 255
Tyr Arg Glu Lys Leu Thr Ile Gly Cys Arg Gln Leu Val Glu Met Glu
            260                 265                 270
Tyr Thr Met Gln Gln Cys Asn Ala Ser Val Tyr Met Glu Ala Lys Asn
        275                 280                 285
Arg Gly Trp Cys Glu Asp Met Leu Asn Tyr Arg Ile  *
    290                 295                 300
```

<210> 1323
<211> 85
<212> PRT
<213> Homo sapiens

<400> 1323

```
Met Thr Glu His Leu Ala Gln Gln Ser Glu Phe Ala Ala Thr Leu Leu
 1               5                  10                  15
Leu Leu Trp Ala Pro Leu Lys Thr Gly Arg Leu Thr Asn Ser Phe Val
            20                  25                  30
Asn Gly Pro Gly Gln His Gly Lys Met Cys Cys Ile Leu Pro Pro Lys
        35                  40                  45
Thr Pro Val Ser Thr Lys Asn Ala Lys Ile Gly Arg Ala Trp Trp Cys
    50                  55                  60
Thr Ser Val Ile Pro Ala Thr Trp Glu Ala Asp Thr Gly Glu Ser Leu
65                  70                  75                  80
Glu Pro Gly Arg  *
            84
```

<210> 1324
<211> 46
<212> PRT
<213> Homo sapiens

<400> 1324
Met Leu His His Ser Gln Leu Ile Phe Val Phe Leu Val Gln Thr Gly
1 5 10 15
Phe His His Val Ala Leu Ser Gly Phe Lys Leu Leu Ala Ser Ser Asn
20 25 30
Leu Pro Thr Leu Asp Pro Lys Val Leu Gly Leu Gln Val *
35 40 45

<210> 1325
<211> 87
<212> PRT
<213> Homo sapiens

<400> 1325
Met Gly Leu Ser Lys Ala Phe Leu Ile Thr Arg Thr Val Phe Leu Ile
1 5 10 15
Ser Ser Leu Ser Phe Tyr Ser Phe Leu Gly Phe Pro Ser Leu Cys Phe
20 25 30
Thr Gly Ser Cys Met Leu Ser Thr Leu Phe Ile Arg Ala Leu Ser Ile
35 40 45
Leu Val Ile Ile Val Leu Asn Ser Arg Ser Asp Lys Ser Asn Thr Pro
50 55 60
Ala Ile Ser Glu Ser Gly Ser Asp Ala Cys Ser Phe Ser Ser Asn Phe
65 70 75 80
Val Phe Cys Leu Leu Val *
85 86

<210> 1326
<211> 69
<212> PRT
<213> Homo sapiens

<400> 1326
Met Ser Leu Phe Leu Phe Phe Leu Met Phe Gln Val Leu Ser Glu Val
1 5 10 15
Ser Trp Gly Gly Val Gly Ser Val Ser Asn Gln Gly Leu Glu His His
20 25 30
Glu Ile Val Thr Pro Asp Leu Gln Ser Leu Ala Gly Gly Trp Thr Gly
35 40 45
Gly Arg Glu Arg Gly Phe Leu Phe Thr Phe Asn Ile Phe Leu Gln Lys
50 55 60
Lys Gln Thr Ile *
65 68

<210> 1327
<211> 103
<212> PRT
<213> Homo sapiens

<221> misc_feature

<222> (1)...(103)
<223> Xaa = any amino acid or nothing

<400> 1327
Met Val Gly Phe Gly Thr Asn Arg Arg Ala Gly Arg Leu Pro Ser Leu
1 5 10 15
Val Leu Val Val Leu Leu Val Val Ile Val Val Leu Ala Phe Asn Tyr
20 25 30
Trp Ser Ile Ser Ser Arg His Val Leu Leu Glu Glu Glu Val Ala Glu
35 40 45
Leu Gln Gly Arg Val Gln Arg Ala Glu Val Ala Leu Trp Arg Val Gly
50 55 60
Gly Arg Asn Cys Asp Leu Leu Leu Val Val Gly Thr Arg Ser Arg Arg
65 70 75 80
Ile Glu Glu Arg Gly Ala Asp Tyr Ser Arg Leu Ser Arg Arg Leu Gln
85 90 95
Xaa Lys Glu Gly Leu Val Asn
100 103

<210> 1328
<211> 52
<212> PRT
<213> Homo sapiens

<400> 1328
Met Arg Ala Arg Pro Ala Cys Thr Ala Thr Phe Pro Ser Phe His Leu
1 5 10 15
Ala Leu Asp Ser Ser Tyr Leu Pro Cys Cys Lys Gly Lys Ala Thr Phe
20 25 30
Ile Pro Lys Ser Arg Ile Tyr Leu Gln Glu Ala Lys Gly Ser Gly Glu
35 40 45
Pro Leu Gly *
50 51

<210> 1329
<211> 204
<212> PRT
<213> Homo sapiens

<400> 1329
Met Cys Thr Arg Asn Leu Ala Leu Leu Phe Ala Pro Ser Val Phe Gln
1 5 10 15
Thr Asp Gly Arg Gly Glu His Glu Val Arg Val Leu Gln Glu Leu Ile
20 25 30
Asp Gly Tyr Ile Ser Val Phe Asp Ile Asp Ser Asp Gln Val Ala Gln
35 40 45
Ile Asp Leu Glu Val Ser Leu Ile Thr Thr Trp Lys Asp Val Gln Leu
50 55 60
Ser Gln Ala Gly Asp Leu Ile Met Glu Val Tyr Ile Glu Gln Gln Leu
65 70 75 80
Pro Asp Asn Cys Val Thr Leu Lys Val Ser Pro Thr Leu Thr Ala Glu
85 90 95

Glu Leu Thr Asn Gln Val Leu Glu Met Arg Gly Thr Ala Ala Gly Met
100 105 110
Asp Leu Trp Val Thr Phe Glu Ile Arg Glu His Gly Glu Leu Glu Arg
115 120 125
Pro Leu His Pro Lys Glu Lys Val Leu Glu Gln Ala Leu Gln Trp Cys
130 135 140
Gln Leu Pro Glu Pro Cys Ser Ala Ser Leu Leu Leu Lys Lys Val Pro
145 150 155 160
Leu Ala Gln Ala Gly Cys Leu Phe Thr Gly Ile Arg Arg Glu Ser Pro
165 170 175
Arg Val Gly Leu Phe Ala Val Phe Val Arg Ser His Leu Ala Cys Trp
180 185 190
Gly Ser Arg Phe Gln Glu Arg Phe Phe Leu Val Ala
195 200 204

<210> 1330
<211> 199
<212> PRT
<213> Homo sapiens

<400> 1330
Met Pro Val Pro Ala Leu Cys Leu Leu Trp Ala Leu Ala Met Val Thr
1 5 10 15
Arg Pro Ala Ser Ala Ala Pro Met Gly Gly Pro Glu Leu Ala Gln His
20 25 30
Glu Glu Leu Thr Leu Leu Phe His Gly Thr Leu Gln Leu Gly Gln Ala
35 40 45
Leu Asn Gly Val Tyr Arg Thr Thr Glu Gly Arg Leu Thr Lys Ala Arg
50 55 60
Asn Ser Leu Gly Leu Tyr Gly Arg Thr Ile Glu Leu Leu Gly Gln Glu
65 70 75 80
Val Ser Arg Gly Arg Asp Ala Ala Gln Glu Leu Arg Ala Ser Leu Leu
85 90 95
Glu Thr Gln Met Glu Glu Asp Ile Leu Gln Leu Gln Ala Glu Ala Thr
100 105 110
Ala Glu Val Leu Gly Glu Val Ala Gln Ala Gln Lys Val Leu Arg Asp
115 120 125
Ser Val Gln Arg Leu Glu Val Gln Leu Arg Ser Ala Trp Leu Gly Pro
130 135 140
Ala Tyr Arg Glu Phe Glu Val Leu Lys Ala His Ala Asp Lys Gln Ser
145 150 155 160
His Ile Leu Trp Ala Leu Thr Gly His Val Gln Arg Gln Arg Arg Glu
165 170 175
Met Val Ala Gln Gln His Arg Leu Arg Gln Ile Gln Glu Arg Leu His
180 185 190
Thr Ala Ala Leu Pro Ala *
195 198

<210> 1331
<211> 81
<212> PRT
<213> Homo sapiens

<400> 1331
Met Ala Arg Pro Ser Ala Phe Pro Ile Gly Val Cys Leu Thr Leu Pro
1 5 10 15
Met Ala Trp Ile Ser Pro Gly Leu Ala Val Pro Ser Cys Pro Gln Tyr
20 25 30
Ile Leu Gln Ala Gln Gly Cys Ile Leu Asp Met Lys Thr Arg Gly Ser
35 40 45
His Gly Glu Ser Ala Val Pro Gly Ala His Gly Ser Arg Pro Phe His
50 55 60
Pro Leu Ala Glu Pro Asn Pro Pro Arg Gln Lys Leu Thr Pro Cys Thr
65 70 75 80
*

<210> 1332
<211> 73
<212> PRT
<213> Homo sapiens

<221> misc_feature
<222> (1)...(73)
<223> Xaa = any amino acid or nothing

<400> 1332
Met Thr Ile Ile Leu Gln Ile Glu Thr Val Ile Phe Leu Leu Tyr Leu
1 5 10 15
Ala Pro Asp Thr Val Arg Pro Leu Thr Ile Ile Thr Gly Met Ala Gly
20 25 30
Ile Val Lys Gln Gln Ile Asp Ser His Ile Thr Asp Pro Asp Gln Gln
35 40 45
Asn Asn Gly Leu Ser Leu Ser Gly Pro Pro Pro Ala Pro Asp Pro Leu
50 55 60
Asp Xaa Leu Val Pro Thr Leu Trp Gly
65 70 73

<210> 1333
<211> 52
<212> PRT
<213> Homo sapiens

<400> 1333
Met Leu Val Tyr Ile Leu Trp Asn Met Tyr Phe Asn Val Cys Ile Val
1 5 10 15
Pro Gly Val Ile Lys Ser Lys Thr Gly Thr Gln Asp Leu Ser Gly Leu
20 25 30
Trp Pro Leu Gly Thr Phe Pro Leu Ile Thr Phe Leu Pro Thr Trp Leu
35 40 45
Ser Tyr Gly *
50 51

<210> 1334

<211> 65
<212> PRT
<213> Homo sapiens

<400> 1334
Met Ile Leu Phe Gln Leu Pro Ser Asn Val Phe Val Leu Leu Met Phe
1               5                   10                  15
Leu Phe Leu Phe Glu Phe Phe Leu Thr Leu Val Pro Met Trp Ala Phe
            20                  25                  30
Pro Gly Asp Lys Thr Phe Val Ser Pro Ala Ser Ser Leu Ser Phe Leu
        35                  40                  45
Asp Leu Ser Phe Leu Leu Phe Cys Asn Ser Val Ser Ile Gly Lys Gln
    50                  55                  60                  64
*

<210> 1335
<211> 112
<212> PRT
<213> Homo sapiens

<400> 1335
Met Leu His Pro Glu Thr Ser Pro Gly Arg Gly His Leu Leu Ala Val
1               5                   10                  15
Leu Leu Ala Leu Leu Gly Thr Ala Trp Ala Glu Val Trp Pro Pro Gln
            20                  25                  30
Leu Gln Glu Gln Ala Pro Met Ala Gly Ala Leu Asn Arg Lys Glu Ser
        35                  40                  45
Phe Leu Leu Leu Ser Leu His Asn Arg Leu Arg Ser Trp Val Gln Pro
    50                  55                  60
Pro Ala Ala Asp Met Arg Arg Leu Asp Trp Ser Asp Ser Leu Ala Gln
65                  70                  75                  80
Leu Ala Gln Ala Arg Ala Ala Leu Cys Gly Ile Pro Thr Pro Ser Leu
                85                  90                  95
Ala Ser Gly Leu Trp Arg Thr Leu Gln Val Gly Trp Asn Met Gln Leu
            100                 105                 110     112

<210> 1336
<211> 105
<212> PRT
<213> Homo sapiens

<400> 1336
Met Thr Gly Asn Leu Cys Phe Phe Ser Ile Lys Gly Tyr Leu Leu Thr
1               5                   10                  15
Ser Glu Ile Leu Met Ile Tyr Leu Thr Leu Glu Phe Cys Ile Leu Arg
            20                  25                  30
Gly Lys His Leu Asn Val Ser Phe Lys Ala Gly Asp Thr Phe Ile Leu
        35                  40                  45
Tyr Leu Gly Ser Leu Gly Phe Glu Glu Glu Gly Gly Pro Glu Ile Leu

```
        50                  55                  60
Lys Asp Cys Met Gly Gly Leu Ser Ser Pro Pro Leu Trp Lys Ala Glu
65                  70                  75                  80
Ala Gly Cys Ile Ile Trp Gly Leu Gly Val Trp Asp His Pro Trp Ala
                85                  90                  95
Thr Thr Arg His Pro Leu Leu Cys  *
            100             104


<210> 1337
<211> 57
<212> PRT
<213> Homo sapiens


<400> 1337
Met Tyr Val Leu Ser Ser Ala His Leu Cys Phe Leu Cys Leu Gln Cys
 1               5                  10                  15
Ser Ser Leu Glu Val Tyr Leu Ile Ser Ser Leu Thr Ser Phe Arg Ser
            20                  25                  30
Val Leu Asn Cys Tyr Pro Pro Glu Arg Ser Ser Leu Thr Ile Gln Tyr
        35                  40                  45
Gln Ile Leu Leu Leu Leu Leu Gln  *
    50                  55  56


<210> 1338
<211> 59
<212> PRT
<213> Homo sapiens


<400> 1338
Met Arg Ile Ile Ser Leu Thr Leu Met Leu Leu Glu Leu Phe Asp Ser
 1               5                  10                  15
Glu Asp Pro Arg Gln Arg Glu Tyr Leu Lys Asn Ile Leu His Arg Leu
            20                  25                  30
Tyr Gly Arg Met Leu Gly Leu Arg Pro Tyr Ile His Lys Gln Ser Lys
        35                  40                  45
His Ile Phe Leu Arg Met Ile Tyr Glu Phe  *
    50                  55          58


<210> 1339
<211> 50
<212> PRT
<213> Homo sapiens


<400> 1339
Met Ile Lys Leu Ala Ile Trp Ser Ile Ile Ile Gly Leu Arg Leu Thr
 1               5                  10                  15
Ile Leu Phe Cys Ile Glu Thr Arg Glu Ser Asp Ile Cys Lys Ile Leu
            20                  25                  30
Gln Tyr Thr Glu Ser Thr Ile Phe Trp Arg Phe Phe Pro Val Tyr Arg
        35                  40                  45
```

Tyr *
49

<210> 1340
<211> 81
<212> PRT
<213> Homo sapiens

<400> 1340
Met Pro Leu Ala Cys Thr Gly Leu Asn Thr Gln Arg Phe Ser Tyr Leu
1 5 10 15
Arg Asp Leu Phe Leu Pro Trp Gly Leu Cys Ile Leu Tyr Ser Ile Leu
20 25 30
Ser Ala Ile Phe Pro Asp Leu Ser Ser Ser Ala Lys Leu Pro Ser Leu
35 40 45
His Ile Ala Phe Phe Thr Leu Phe Lys Val Thr Lys Gly Thr Ser Pro
50 55 60
Lys Ala Thr Asp Val Pro Val Ala Cys Phe Ile Asn His Asn Arg Thr
65 70 75 80
*

<210> 1341
<211> 60
<212> PRT
<213> Homo sapiens

<400> 1341
Met Phe Glu Ile His Arg Ala His Gly Val Phe Leu Leu Leu Ser Ile
1 5 10 15
Gln Leu Thr Thr Ser Leu Lys Arg Lys Ser Gly Glu Gly Asp Arg Glu
20 25 30
Ser Pro Ala Ser Trp Phe Ser Pro Phe Ser Gln Met Phe Phe Leu Ile
35 40 45
Asn Thr Ile Leu Leu Pro Phe Lys Ile Pro Ile *
50 55 59

<210> 1342
<211> 49
<212> PRT
<213> Homo sapiens

<400> 1342
Met Leu Ser Leu Phe Ile Phe Leu Arg Phe Leu Pro Leu Gly Phe Cys
1 5 10 15
Trp Lys Glu Leu His Pro Glu Ala Glu Gln Ser Glu Lys Val Asp Phe
20 25 30
Arg Lys Pro Trp Tyr Leu Thr Gly His Ala Ala Ser Leu Gly Ala Asp
35 40 45 48
*

<210> 1343
<211> 70
<212> PRT
<213> Homo sapiens

<400> 1343
```
Met Arg Leu Ala Val Ser Cys Ile Thr Ser Phe Leu Met Leu Ser Leu
 1               5                  10                  15
Leu Leu Phe Met Ala His Arg Leu Arg Gln Arg Arg Arg Glu Arg Ile
            20                  25                  30
Glu Ser Leu Ile Gly Ala Asn Leu His His Phe Asn Leu Gly Arg Arg
        35                  40                  45
Ile Pro Gly Phe Asp Tyr Gly Pro Asp Gly Phe Gly Thr Gly Leu Thr
    50                  55                  60
Pro Leu Ala Phe Phe  *
65              69
```

<210> 1344
<211> 99
<212> PRT
<213> Homo sapiens

<400> 1344
```
Met Phe Leu Ser Leu Ser Leu Thr Leu Cys Leu Cys Phe Ser Phe Phe
 1               5                  10                  15
Cys Leu Tyr Leu Ser Leu Ala Leu Tyr Leu Gly Ser Phe Phe Cys Leu
            20                  25                  30
Pro Phe His Val Ser Val Phe Leu Cys Leu Phe Pro Ser Val Leu Phe
        35                  40                  45
Leu Ser Val Ala Leu Gly Ser Pro Glu Asn His Ile Ser Trp Arg Lys
    50                  55                  60
Val Gly Glu Glu Leu Lys Leu Ala Ser His Arg Asn Phe Cys Ser Leu
65                  70                  75                  80
Met Gln Lys Met Arg Ser Asn Lys Pro Ser Pro Ser Arg Pro Arg Gly
                85                  90                  95
Trp Ala  *
    98
```

<210> 1345
<211> 112
<212> PRT
<213> Homo sapiens

<400> 1345
```
Met Lys Val Leu Trp Ala Gly Val Leu Gly Thr Phe Leu Ala Gly Cys
 1               5                  10                  15
Gln Ala Lys Val Glu Gln Ala Val Glu Thr Glu Pro Glu Pro Glu Leu
            20                  25                  30
```

Cys Gln Gln Thr Glu Trp Lys Ser Gly Gln Arg Trp Glu Leu Glu Leu
35 40 45
Gly Arg Phe Trp Asp Tyr Leu Arg Trp Glu Gln Thr Leu Ser Glu Gln
50 55 60
Val Gln Glu Glu Leu Val Ser Ser Gln Val Thr Gln Glu Leu Lys Ala
65 70 75 80
Leu Met Asp Glu Thr Met Lys Glu Met Lys Ala Tyr Lys Ser Asp Leu
85 90 95
Glu Glu Gln Leu Thr Pro Val Ala Gly Arg Arg Trp His Gly Cys Thr
100 105 110 112

<210> 1346
<211> 360
<212> PRT
<213> Homo sapiens

<400> 1346
Met Leu Phe Val Pro Val Thr Leu Cys Met Ile Val Val Val Ala Thr
1 5 10 15
Ile Lys Ser Val Arg Phe Tyr Thr Glu Lys Asn Gly Gln Leu Ile Tyr
20 25 30
Thr Pro Phe Thr Glu Asp Thr Pro Ser Val Gly Gln Arg Leu Leu Asn
35 40 45
Ser Val Leu Asn Thr Leu Ile Met Ile Ser Val Ile Val Val Met Thr
50 55 60
Ile Phe Leu Val Val Leu Tyr Lys Tyr Arg Cys Tyr Lys Phe Ile His
65 70 75 80
Gly Trp Leu Ile Met Ser Ser Leu Met Leu Leu Phe Leu Phe Thr Tyr
85 90 95
Ile Tyr Leu Gly Glu Val Leu Lys Thr Tyr Asn Val Ala Met Asp Tyr
100 105 110
Pro Thr Leu Leu Leu Thr Val Trp Asn Phe Gly Ala Val Gly Met Val
115 120 125
Cys Ile His Trp Lys Gly Pro Leu Val Leu Gln Gln Ala Tyr Leu Ile
130 135 140
Met Ile Ser Ala Leu Met Ala Leu Val Phe Ile Lys Tyr Leu Pro Glu
145 150 155 160
Trp Ser Ala Trp Val Ile Leu Gly Ala Ile Ser Val Tyr Asp Leu Val
165 170 175
Ala Val Leu Cys Pro Lys Gly Pro Leu Arg Met Leu Val Glu Thr Ala
180 185 190
Gln Glu Arg Asn Glu Pro Ile Phe Pro Ala Leu Ile Tyr Ser Ser Ala
195 200 205
Met Val Trp Thr Val Gly Met Ala Lys Leu Asp Pro Ser Ser Gln Gly
210 215 220
Ala Leu Gln Leu Pro Tyr Asp Pro Glu Met Glu Glu Asp Ser Tyr Asp
225 230 235 240
Ser Phe Gly Glu Pro Ser Tyr Pro Glu Val Phe Glu Pro Pro Leu Thr
245 250 255
Gly Tyr Pro Gly Glu Glu Leu Glu Glu Glu Glu Glu Arg Gly Val Lys
260 265 270
Leu Gly Leu Gly Asp Phe Ile Phe Tyr Ser Val Leu Val Gly Lys Ala
275 280 285
Ala Ala Thr Gly Ser Gly Asp Trp Asn Thr Thr Leu Ala Cys Phe Val

```
    290                 295                 300
Ala Ile Leu Ile Gly Leu Cys Leu Thr Leu Leu Leu Leu Ala Val Phe
305                 310                 315                 320
Lys Lys Ala Leu Pro Ala Leu Pro Ile Ser Ile Thr Phe Gly Leu Ile
                325                 330                 335
Phe Tyr Phe Ser Thr Asp Asn Leu Val Arg Pro Phe Met Asp Thr Leu
            340                 345                 350
Ala Ser His Gln Leu Tyr Ile  *
        355             359
```

```
<210> 1347
<211> 84
<212> PRT
<213> Homo sapiens
```

```
<400> 1347
Met Ile Leu Ser Leu Tyr Tyr Lys Leu Phe Gly Lys Leu Ala Val Ala
 1               5                  10                  15
Thr Ile Glu Ile Leu His Cys Leu Cys Tyr Ile Glu Phe Val Ile Ile
            20                  25                  30
Phe Lys Gly Phe Lys Lys Ile Pro Ile Cys Phe Phe Ser Phe Leu Phe
        35                  40                  45
Ser Phe Val Pro His His Leu Asn Tyr Leu Gly Lys Tyr His Ser Ser
    50                  55                  60
Lys Phe Glu Tyr Cys Leu Ser Asn Lys Lys Lys Cys Glu Arg Tyr Glu
65                  70                  75                  80
Glu Glu Arg  *
        83
```

```
<210> 1348
<211> 65
<212> PRT
<213> Homo sapiens
```

```
<400> 1348
Met Val His Leu Leu Leu Val Phe Trp Ser Gly Pro His Asn Leu Gly
 1               5                  10                  15
Arg Phe Gln Pro Met Lys Leu Phe Ala Ile Cys Leu Asn Gln Ser Gly
            20                  25                  30
Tyr Ile Ile Ala Phe Phe Val Leu Tyr Thr Asn Arg Met Tyr Ser Ile
        35                  40                  45
Ile Asn Ile Ile Leu Asn Leu Phe Tyr Pro Val Tyr Tyr Cys Lys Ile
    50                  55                  60                  64
 *
```

```
<210> 1349
<211> 58
<212> PRT
<213> Homo sapiens
```

<400> 1349
Met Pro Ser Pro Ser Gly Leu Trp Arg Ile Leu Leu Leu Val Leu Gly
1 5 10 15
Ser Val Leu Ser Gly Ser Ala Arg Ala Ala Ala Pro Leu Arg Val Leu
20 25 30
Arg Gln Thr Ala Leu Cys Cys Ala Thr Glu Ala Leu Val Ala Val Pro
35 40 45
Glu Gly Ile Pro Thr Glu Thr Arg Leu *
50 55 57

<210> 1350
<211> 60
<212> PRT
<213> Homo sapiens

<221> misc_feature
<222> (1)...(60)
<223> Xaa = any amino acid or nothing

<400> 1350
Met Gly Ile Gly Cys Trp Arg Asn Pro Leu Val Leu Leu Met Ala Leu
1 5 10 15
Ala Cys Gln Ala Ser Trp Gly Leu Ser Lys Gly Gly Arg Val Leu Pro
20 25 30
Asn Leu Cys Pro Lys Lys Met Phe Xaa Thr Leu Phe Phe Phe Asn Ser
35 40 45
Gln Arg Gly Arg Gly Pro Pro Phe Trp Ala Gly Gly
50 55 60

<210> 1351
<211> 56
<212> PRT
<213> Homo sapiens

<400> 1351
Met Leu Leu Ala Leu Pro Leu Ala Ala Pro Ser Cys Pro Met Leu Cys
1 5 10 15
Thr Cys Tyr Ser Ser Pro Pro Thr Val Ser Cys Gln Ala Asn Asn Phe
20 25 30
Ser Ser Val Pro Leu Ser Leu Pro Pro Ser Thr Gln Arg Leu Phe Leu
35 40 45
Gln Asn Asn Leu Ile Arg Thr Leu
50 55 56

<210> 1352
<211> 701
<212> PRT
<213> Homo sapiens

```
        <400> 1352
Met Glu Pro Leu Cys Pro Leu Leu Leu Val Gly Phe Ser Leu Pro Leu
 1               5                  10                  15
Ala Arg Ala Leu Arg Gly Asn Glu Thr Thr Ala Asp Ser Asn Glu Thr
            20                  25                  30
Thr Thr Thr Ser Gly Pro Pro Asp Pro Gly Ala Ser Gln Pro Leu Leu
        35                  40                  45
Ala Trp Leu Leu Leu Pro Leu Leu Leu Leu Leu Leu Val Leu Leu Leu
    50                  55                  60
Ala Ala Tyr Phe Phe Arg Phe Arg Lys Gln Arg Lys Ala Val Val Ser
65                  70                  75                  80
Thr Ser Asp Lys Lys Met Pro Asn Gly Ile Leu Glu Glu Gln Glu Gln
                85                  90                  95
Gln Arg Val Met Leu Leu Ser Arg Ser Pro Ser Gly Pro Lys Lys Tyr
            100                 105                 110
Phe Pro Ile Pro Val Glu His Leu Glu Glu Glu Ile Arg Ile Arg Ser
        115                 120                 125
Ala Asp Asp Cys Lys Gln Phe Arg Glu Glu Phe Asn Ser Leu Pro Ser
    130                 135                 140
Gly His Ile Gln Gly Thr Phe Glu Leu Ala Asn Lys Glu Glu Asn Arg
145                 150                 155                 160
Glu Lys Asn Arg Tyr Pro Asn Ile Leu Pro Asn Asp His Ser Arg Val
                165                 170                 175
Ile Leu Ser Gln Leu Asp Gly Ile Pro Cys Ser Asp Tyr Ile Asn Ala
            180                 185                 190
Ser Tyr Ile Asp Gly Tyr Lys Glu Lys Asn Lys Phe Ile Ala Ala Gln
        195                 200                 205
Gly Pro Lys Gln Glu Thr Val Asn Asp Phe Trp Arg Met Val Trp Glu
    210                 215                 220
Gln Lys Ser Ala Thr Ile Val Met Leu Thr Asn Leu Lys Glu Arg Lys
225                 230                 235                 240
Glu Glu Lys Cys His Gln Tyr Trp Pro Asp Gln Gly Cys Trp Thr Tyr
                245                 250                 255
Gly Asn Ile Arg Val Cys Val Glu Asp Cys Val Val Leu Val Asp Tyr
            260                 265                 270
Thr Ile Arg Lys Phe Cys Ile Gln Pro Gln Leu Pro Asp Gly Cys Lys
        275                 280                 285
Ala Pro Arg Leu Val Ser Gln Leu His Phe Thr Ser Trp Pro Asp Phe
    290                 295                 300
Gly Val Pro Phe Thr Pro Ile Gly Met Leu Lys Phe Leu Lys Lys Val
305                 310                 315                 320
Lys Thr Leu Asn Pro Val His Ala Gly Pro Ile Val Val His Cys Ser
                325                 330                 335
Ala Gly Val Gly Arg Thr Gly Thr Phe Ile Val Ile Asp Ala Met Met
            340                 345                 350
Ala Met Met His Ala Glu Gln Lys Val Asp Val Phe Glu Phe Val Ser
        355                 360                 365
Arg Ile Arg Asn Gln Arg Pro Gln Met Val Gln Thr Asp Met Gln Tyr
    370                 375                 380
Thr Phe Ile Tyr Gln Ala Leu Leu Glu Tyr Tyr Leu Tyr Gly Asp Thr
385                 390                 395                 400
Glu Leu Asp Val Ser Ser Leu Glu Lys His Leu Gln Thr Met His Gly
                405                 410                 415
Thr Thr Thr His Phe Asp Lys Ile Gly Leu Glu Glu Glu Phe Arg Lys
            420                 425                 430
Leu Thr Asn Val Arg Ile Met Lys Glu Asn Met Arg Thr Gly Asn Leu
        435                 440                 445
Pro Ala Asn Met Lys Lys Ala Arg Val Ile Gln Ile Ile Pro Tyr Asp
    450                 455                 460
```

Phe Asn Arg Val Ile Leu Ser Met Lys Arg Gly Gln Glu Tyr Thr Asp
465 470 475 480
Tyr Ile Asn Ala Ser Phe Ile Asp Gly Tyr Arg Gln Lys Asp Tyr Phe
485 490 495
Ile Ala Thr Gln Gly Pro Leu Ala His Thr Val Glu Asp Phe Trp Arg
500 505 510
Met Ile Trp Glu Trp Lys Ser His Thr Ile Val Met Leu Thr Glu Val
515 520 525
Gln Glu Arg Glu Gln Asp Lys Cys Tyr Gln Tyr Trp Pro Thr Glu Gly
530 535 540
Ser Val Thr His Gly Glu Ile Thr Ile Glu Ile Lys Asn Asp Thr Leu
545 550 555 560
Ser Glu Ala Ile Ser Ile Arg Asp Phe Leu Val Thr Leu Asn Gln Pro
565 570 575
Gln Ala Arg Gln Glu Glu Gln Val Arg Val Val Arg Gln Phe His Phe
580 585 590
His Gly Trp Pro Glu Ile Gly Ile Pro Ala Glu Gly Lys Gly Met Ile
595 600 605
Asp Leu Ile Ala Ala Val Gln Lys Gln Gln Gln Gln Thr Gly Asn His
610 615 620
Pro Ile Thr Val His Cys Ser Ala Gly Ala Gly Arg Thr Gly Thr Phe
625 630 635 640
Ile Ala Leu Ser Asn Ile Leu Glu Arg Val Lys Ala Glu Gly Leu Leu
645 650 655
Asp Val Phe Gln Ala Val Lys Ser Leu Arg Leu Gln Arg Pro His Met
660 665 670
Val Gln Thr Leu Glu Gln Tyr Glu Phe Cys Tyr Lys Val Val Gln Asp
675 680 685
Phe Ile Asp Ile Phe Ser Asp Tyr Ala Asn Phe Lys *
690 695 700

<210> 1353
<211> 49
<212> PRT
<213> Homo sapiens

<400> 1353
Met Ala Phe Leu Tyr His Val Ala Tyr Val Leu Val Cys Met Leu Gly
1 5 10 15
Leu Phe Cys His Glu Phe Phe Tyr Ser Phe Leu Leu Phe Glu Ser Val
20 25 30
Tyr Arg His Gln Thr Leu Leu Asn Asp Ile Pro Cys Val Lys Leu Met
35 40 45 48
*

<210> 1354
<211> 58
<212> PRT
<213> Homo sapiens

<400> 1354
Met Ser Val Cys Lys Tyr Thr Val Tyr Gly Phe Phe Ile Phe Ala Phe

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 1               5                   10                  15
Phe Tyr Phe Thr Lys Asp Asn Ile Pro Tyr Leu Lys Val Ser Leu Gln
            20                  25                  30
Ala Phe Cys Gly Phe Gln Asn Ile Ser Trp Asn Lys Tyr Thr Leu Leu
        35                  40                  45
Phe Tyr Tyr Ser Pro Leu Thr Ile Ile  *
    50                  55      57
```

<210> 1355
<211> 4261
<212> PRT
<213> Homo sapiens

<400> 1355

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Met Leu Ser Ala Ile Leu Leu Leu Leu Gln Leu Trp Asp Ser Gly Ala
 1               5                   10                  15
Gln Glu Thr Asp Asn Glu Arg Ser Ala Gln Gly Thr Ser Ala Pro Leu
            20                  25                  30
Leu Pro Leu Leu Gln Arg Phe Gln Ser Ile Ile Cys Arg Lys Asp Ala
        35                  40                  45
Pro His Ser Glu Gly Asp Met His Leu Leu Ser Gly Pro Leu Ser Pro
    50                  55                  60
Asn Glu Ser Phe Leu Arg Tyr Leu Thr Leu Pro Gln Asp Asn Glu Leu
65                  70                  75                  80
Ala Ile Asp Leu Arg Gln Thr Ala Val Val Val Met Ala His Leu Asp
                85                  90                  95
Arg Leu Ala Thr Pro Cys Met Pro Pro Leu Cys Ser Ser Pro Thr Ser
            100                 105                 110
His Lys Gly Ser Leu Gln Glu Val Ile Gly Trp Gly Leu Ile Gly Trp
        115                 120                 125
Lys Tyr Tyr Ala Asn Val Ile Gly Pro Ile Gln Cys Glu Gly Leu Ala
    130                 135                 140
Asn Leu Gly Val Thr Gln Ile Ala Cys Ala Glu Lys Arg Phe Leu Ile
145                 150                 155                 160
Leu Ser Arg Asn Gly Arg Val Tyr Thr Gln Ala Tyr Asn Ser Asp Thr
                165                 170                 175
Leu Ala Pro Gln Leu Val Gln Gly Leu Ala Ser Arg Asn Ile Val Lys
            180                 185                 190
Ile Ala Ala His Ser Asp Gly His His Tyr Leu Ala Leu Ala Ala Thr
        195                 200                 205
Gly Glu Val Tyr Ser Trp Gly Cys Gly Asp Gly Gly Arg Leu Gly His
    210                 215                 220
Gly Asp Thr Val Pro Leu Glu Glu Pro Lys Val Ile Ser Ala Phe Ser
225                 230                 235                 240
Gly Lys Gln Ala Gly Lys His Val Val His Ile Ala Cys Gly Ser Thr
                245                 250                 255
Tyr Ser Ala Ala Ile Thr Ala Glu Gly Glu Leu Tyr Thr Trp Gly Arg
            260                 265                 270
Gly Asn Tyr Gly Arg Leu Gly His Gly Ser Ser Glu Asp Glu Ala Ile
        275                 280                 285
Pro Met Leu Val Ala Gly Leu Lys Gly Leu Lys Val Ile Asp Val Ala
    290                 295                 300
Cys Gly Ser Gly Asp Ala Gln Thr Leu Ala Val Thr Glu Asn Gly Gln
305                 310                 315                 320
Val Trp Ser Trp Gly Asp Gly Asp Tyr Gly Lys Leu Gly Arg Gly Gly
                325                 330                 335
```

Ser Asp Gly Cys Lys Thr Pro Lys Leu Ile Glu Lys Leu Gln Asp Leu
340 345 350
Asp Val Val Lys Val Arg Cys Gly Ser Gln Phe Ser Ile Ala Leu Thr
355 360 365
Lys Asp Gly Gln Val Tyr Ser Trp Gly Lys Gly Asp Asn Gln Arg Leu
370 375 380
Gly His Gly Thr Glu Glu His Val Arg Tyr Pro Lys Leu Leu Glu Gly
385 390 395 400
Leu Gln Gly Lys Lys Val Ile Asp Val Ala Ala Gly Ser Thr His Cys
405 410 415
Leu Ala Leu Thr Glu Asp Ser Glu Val His Ser Trp Gly Ser Asn Asp
420 425 430
Gln Cys Gln His Phe Asp Thr Leu Arg Val Thr Lys Pro Glu Pro Ala
435 440 445
Ala Leu Pro Gly Leu Asp Thr Lys His Ile Val Gly Ile Ala Cys Gly
450 455 460
Pro Ala Gln Ser Phe Ala Trp Ser Ser Cys Ser Glu Trp Ser Ile Gly
465 470 475 480
Leu Arg Val Pro Phe Val Val Asp Ile Cys Ser Met Thr Phe Glu Gln
485 490 495
Leu Asp Leu Leu Leu Arg Gln Val Ser Glu Gly Met Asp Gly Ser Ala
500 505 510
Asp Trp Pro Pro Pro Gln Glu Lys Glu Cys Val Ala Val Ala Thr Leu
515 520 525
Asn Leu Leu Arg Leu Gln Leu His Ala Ala Ile Ser His Gln Val Asp
530 535 540
Pro Glu Phe Leu Gly Leu Gly Leu Gly Ser Ile Leu Leu Asn Ser Leu
545 550 555 560
Lys Gln Thr Val Val Thr Leu Ala Ser Ser Ala Gly Val Leu Ser Thr
565 570 575
Val Gln Ser Ala Ala Gln Ala Val Leu Gln Ser Gly Trp Ser Val Leu
580 585 590
Leu Pro Thr Ala Glu Glu Arg Ala Arg Ala Leu Ser Ala Leu Leu Pro
595 600 605
Cys Ala Val Ser Gly Asn Glu Val Asn Ile Ser Pro Gly Arg Arg Phe
610 615 620
Met Ile Asp Leu Leu Val Gly Ser Leu Met Ala Asp Gly Gly Leu Glu
625 630 635 640
Ser Ala Leu His Ala Ala Ile Thr Ala Glu Ile Gln Asp Ile Glu Ala
645 650 655
Lys Lys Glu Ala Gln Lys Glu Lys Glu Ile Asp Glu Gln Glu Ala Asn
660 665 670
Ala Ser Thr Phe His Arg Ser Arg Thr Pro Leu Asp Lys Asp Leu Ile
675 680 685
Asn Thr Gly Ile Cys Glu Ser Ser Gly Lys Gln Cys Leu Pro Leu Val
690 695 700
Gln Leu Ile Gln Gln Leu Leu Arg Asn Ile Ala Ser Gln Thr Val Ala
705 710 715 720
Arg Leu Lys Asp Val Ala Arg Arg Ile Ser Ser Cys Leu Asp Phe Glu
725 730 735
Gln His Ser Arg Glu Arg Ser Ala Ser Leu Asp Trp Leu Leu Arg Phe
740 745 750
Gln Arg Leu Leu Ile Ser Lys Leu Tyr Pro Gly Glu Ser Ile Gly Gln
755 760 765
Thr Ser Asp Ile Ser Ser Pro Glu Leu Met Gly Val Gly Ser Leu Leu
770 775 780
Lys Lys Tyr Thr Ala Leu Leu Cys Thr His Ile Gly Asp Ile Leu Pro
785 790 795 800
Val Ala Ala Ser Ile Ala Ser Thr Ser Trp Arg His Phe Ala Glu Val

805 810 815
Ala Tyr Ile Val Glu Gly Asp Phe Thr Gly Val Leu Leu Pro Glu Leu
820 825 830
Val Val Ser Ile Val Leu Leu Leu Ser Lys Asn Ala Asp Leu Met Gln
835 840 845
Glu Ala Gly Ala Val Pro Leu Leu Gly Gly Leu Leu Glu His Leu Asp
850 855 860
Arg Phe Asn His Leu Ala Pro Gly Lys Glu Arg Asp Asp His Glu Glu
865 870 875 880
Leu Ala Trp Pro Gly Ile Met Glu Ser Phe Phe Thr Gly Gln Asn Cys
885 890 895
Arg Asn Asn Glu Glu Val Thr Leu Ile Arg Lys Ala Asp Leu Glu Asn
900 905 910
His Asn Lys Asp Gly Gly Phe Trp Thr Val Ile Asp Gly Lys Val Tyr
915 920 925
Asp Ile Lys Asp Phe Gln Thr Gln Ser Leu Thr Gly Asn Ser Ile Leu
930 935 940
Ala Gln Phe Ala Gly Glu Asp Pro Val Val Ala Leu Glu Ala Ala Leu
945 950 955 960
Gln Phe Glu Asp Thr Arg Glu Ser Met His Ala Phe Cys Val Gly Gln
965 970 975
Tyr Leu Glu Pro Asp Gln Glu Ile Val Thr Ile Pro Asp Leu Gly Ser
980 985 990
Leu Ser Ser Pro Leu Ile Asp Thr Glu Arg Asn Leu Gly Leu Leu Leu
995 1000 1005
Gly Leu His Ala Ser Tyr Leu Ala Met Ser Thr Pro Leu Ser Pro Val
1010 1015 1020
Glu Ile Glu Cys Ala Lys Trp Leu Gln Ser Ser Ile Phe Ser Gly Gly
1025 1030 1035 1040
Leu Gln Thr Ser Gln Ile His Tyr Arg Tyr Asn Glu Glu Lys Asp Glu
1045 1050 1055
Asp His Cys Ser Ser Pro Gly Gly Thr Pro Ala Ser Lys Ser Arg Leu
1060 1065 1070
Cys Ser His Arg Arg Ala Leu Gly Asp His Ser Gln Ala Phe Leu Gln
1075 1080 1085
Ala Ile Ala Asp Asn Asn Ile Gln Asp His Asn Val Lys Asp Phe Leu
1090 1095 1100
Cys Gln Ile Glu Arg Tyr Cys Arg Gln Cys His Leu Thr Thr Pro Ile
1105 1110 1115 1120
Met Phe Pro Pro Glu His Pro Val Glu Glu Val Gly Arg Leu Leu Leu
1125 1130 1135
Cys Cys Leu Leu Lys His Glu Asp Leu Gly His Val Ala Leu Ser Leu
1140 1145 1150
Val His Ala Gly Ala Leu Gly Ile Glu Gln Val Lys His Arg Thr Leu
1155 1160 1165
Pro Lys Ser Val Val Asp Val Cys Arg Val Val Tyr Gln Ala Lys Cys
1170 1175 1180
Ser Leu Ile Lys Thr His Gln Glu Gln Gly Arg Ser Tyr Lys Glu Val
1185 1190 1195 1200
Cys Ala Pro Val Ile Glu Arg Leu Arg Phe Leu Phe Asn Glu Leu Arg
1205 1210 1215
Pro Ala Val Cys Asn Asp Leu Ser Ile Met Ser Lys Phe Lys Leu Leu
1220 1225 1230
Ser Ser Leu Pro Arg Trp Arg Arg Ile Ala Gln Lys Ile Ile Arg Glu
1235 1240 1245
Arg Arg Lys Lys Arg Val Pro Lys Lys Pro Glu Ser Met Asp Asp Glu
1250 1255 1260
Glu Lys Ile Gly Asn Glu Glu Ser Asp Leu Glu Glu Ala Cys Ile Leu
1265 1270 1275 1280

Pro His Ser Pro Ile Asn Val Asp Lys Arg Pro Ile Ala Ile Lys Ser
1285 1290 1295
Pro Lys Asp Lys Trp Gln Pro Leu Leu Ser Thr Val Thr Gly Val His
1300 1305 1310
Lys Tyr Lys Trp Leu Lys Gln Asn Val Gln Gly Leu Tyr Pro Gln Ser
1315 1320 1325
Pro Leu Leu Ser Thr Ile Ala Glu Phe Ala Leu Lys Glu Glu Pro Val
1330 1335 1340
Asp Val Glu Lys Met Arg Lys Cys Leu Leu Lys Gln Leu Glu Arg Ala
1345 1350 1355 1360
Glu Val Arg Leu Glu Gly Ile Asp Thr Ile Leu Lys Leu Ala Ser Lys
1365 1370 1375
Asn Phe Leu Leu Pro Ser Val Gln Tyr Ala Met Phe Cys Gly Trp Gln
1380 1385 1390
Arg Leu Ile Pro Glu Gly Ile Asp Ile Gly Glu Pro Leu Thr Asp Cys
1395 1400 1405
Leu Lys Asp Val Asp Leu Ile Pro Pro Phe Asn Arg Met Leu Leu Glu
1410 1415 1420
Val Thr Phe Gly Lys Leu Tyr Ala Trp Ala Val Gln Asn Ile Arg Asn
1425 1430 1435 1440
Val Leu Met Asp Ala Ser Ala Thr Phe Lys Glu Leu Gly Ile Gln Pro
1445 1450 1455
Val Pro Leu Gln Thr Ile Thr Asn Glu Asn Pro Ser Gly Pro Ser Leu
1460 1465 1470
Gly Thr Ile Pro Gln Ala Arg Phe Leu Leu Val Met Leu Ser Met Leu
1475 1480 1485
Thr Leu Gln His Gly Ala Asn Asn Leu Asp Leu Leu Leu Asn Ser Gly
1490 1495 1500
Met Leu Ala Leu Thr Gln Thr Ala Leu Arg Leu Ile Gly Pro Ser Cys
1505 1510 1515 1520
Asp Asn Val Glu Glu Asp Met Asn Ala Ser Ala Gln Gly Ala Ser Ala
1525 1530 1535
Thr Val Leu Glu Glu Thr Arg Lys Glu Thr Ala Pro Val Gln Leu Pro
1540 1545 1550
Val Ser Gly Pro Glu Leu Ala Ala Met Met Lys Ile Gly Thr Arg Val
1555 1560 1565
Met Arg Gly Val Asp Trp Lys Trp Gly Asp Gln Asp Gly Pro Pro Pro
1570 1575 1580
Gly Leu Gly Arg Val Ile Gly Glu Leu Gly Glu Asp Gly Trp Ile Arg
1585 1590 1595 1600
Val Gln Trp Asp Thr Gly Ser Thr Asn Ser Tyr Arg Met Gly Lys Glu
1605 1610 1615
Gly Lys Tyr Asp Leu Lys Leu Ala Glu Leu Pro Ala Ala Ala Gln Pro
1620 1625 1630
Ser Ala Glu Asp Ser Asp Thr Glu Asp Asp Ser Glu Ala Glu Gln Thr
1635 1640 1645
Glu Arg Asn Ile His Pro Thr Ala Met Met Phe Thr Ser Thr Ile Asn
1650 1655 1660
Leu Leu Gln Thr Leu Cys Leu Ser Ala Gly Val His Ala Glu Ile Met
1665 1670 1675 1680
Gln Ser Glu Ala Thr Lys Thr Leu Cys Gly Leu Leu Arg Met Leu Val
1685 1690 1695
Glu Ser Gly Thr Thr Asp Lys Thr Ser Ser Pro Asn Arg Leu Val Tyr
1700 1705 1710
Arg Glu Gln His Arg Ser Trp Cys Thr Leu Gly Phe Val Arg Ser Ile
1715 1720 1725
Ala Leu Thr Pro Gln Val Cys Gly Ala Leu Ser Ser Pro Gln Trp Ile
1730 1735 1740
Thr Leu Leu Met Lys Val Val Glu Gly His Ala Pro Phe Thr Ala Thr

```
1745                1750                1755                1760
Ser Leu Gln Arg Gln Ile Leu Ala Val His Leu Leu Gln Ala Val Leu
                1765                1770                1775
Pro Ser Trp Asp Lys Thr Glu Arg Ala Arg Asp Met Lys Cys Leu Val
            1780                1785                1790
Glu Lys Leu Phe Asp Phe Leu Gly Ser Leu Leu Thr Thr Cys Ser Ser
        1795                1800                1805
Asp Val Pro Leu Leu Arg Glu Ser Thr Leu Arg Arg Arg Arg Val Arg
    1810                1815                1820
Pro Gln Ala Ser Leu Thr Ala Thr His Ser Ser Thr Leu Ala Glu Glu
1825                1830                1835                1840
Val Val Ala Leu Leu Arg Thr Leu His Ser Leu Thr Gln Trp Asn Gly
                1845                1850                1855
Leu Ile Asn Lys Tyr Ile Asn Ser Gln Leu Arg Ser Ile Thr His Ser
            1860                1865                1870
Phe Val Gly Arg Pro Ser Glu Gly Ala Gln Leu Glu Asp Tyr Phe Pro
        1875                1880                1885
Asp Ser Glu Asn Pro Glu Val Gly Gly Leu Met Ala Val Leu Ala Val
    1890                1895                1900
Ile Gly Gly Ile Asp Gly Arg Leu Arg Leu Gly Gly Gln Val Met His
1905                1910                1915                1920
Asp Glu Phe Gly Glu Gly Thr Val Thr Arg Ile Thr Pro Lys Gly Lys
                1925                1930                1935
Ile Thr Val Gln Phe Ser Asp Met Arg Thr Cys Arg Val Cys Pro Leu
            1940                1945                1950
Asn Gln Leu Lys Pro Leu Pro Ala Val Ala Phe Asn Val Asn Asn Leu
        1955                1960                1965
Pro Phe Thr Glu Pro Met Leu Ser Val Trp Ala Gln Leu Val Asn Leu
    1970                1975                1980
Ala Gly Ser Lys Leu Glu Lys His Lys Ile Lys Lys Ser Thr Lys Gln
1985                1990                1995                2000
Ala Phe Ala Gly Gln Val Asp Leu Asp Leu Leu Arg Cys Gln Gln Leu
                2005                2010                2015
Lys Leu Tyr Ile Leu Lys Ala Gly Arg Ala Leu Leu Ser His Gln Asp
            2020                2025                2030
Lys Leu Arg Gln Ile Leu Ser Gln Pro Ala Val Gln Glu Thr Gly Thr
        2035                2040                2045
Val His Thr Asp Asp Gly Ala Val Val Ser Pro Asp Leu Gly Asp Met
    2050                2055                2060
Ser Pro Glu Gly Pro Gln Pro Pro Met Ile Leu Leu Gln Gln Leu Leu
2065                2070                2075                2080
Ala Ser Ala Thr Gln Pro Ser Pro Val Lys Ala Ile Phe Asp Lys Gln
                2085                2090                2095
Glu Leu Glu Ala Ala Ala Leu Ala Val Cys Gln Cys Leu Ala Val Glu
            2100                2105                2110
Ser Thr His Pro Ser Ser Pro Gly Phe Glu Asp Cys Ser Ser Ser Glu
        2115                2120                2125
Ala Thr Thr Pro Val Ala Val Gln His Ile His Pro Ala Arg Val Lys
    2130                2135                2140
Arg Arg Lys Gln Ser Pro Val Pro Ala Leu Pro Ile Val Val Gln Leu
2145                2150                2155                2160
Met Glu Met Gly Phe Ser Arg Arg Asn Ile Glu Phe Ala Leu Lys Ser
                2165                2170                2175
Leu Thr Gly Ala Ser Gly Asn Ala Ser Ser Leu Pro Gly Val Glu Ala
            2180                2185                2190
Leu Val Gly Trp Leu Leu Asp His Ser Asp Ile Gln Val Thr Glu Leu
        2195                2200                2205
Ser Asp Ala Asp Thr Val Ser Asp Glu Tyr Ser Asp Glu Glu Val Val
    2210                2215                2220
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Glu Asp Val Asp Asp Ala Ala Tyr Ser Met Ser Thr Gly Ala Val Val
2225 2230 2235 2240
Thr Glu Ser Gln Thr Tyr Lys Lys Arg Ala Asp Phe Leu Ser Asn Asp
2245 2250 2255
Asp Tyr Ala Val Tyr Val Arg Glu Asn Ile Gln Val Gly Met Met Val
2260 2265 2270
Arg Cys Cys Arg Ala Tyr Glu Glu Val Cys Glu Gly Asp Val Gly Lys
2275 2280 2285
Val Ile Lys Leu Asp Arg Asp Gly Leu His Asp Leu Asn Val Gln Cys
2290 2295 2300
Asp Trp Gln Gln Lys Gly Gly Thr Tyr Trp Val Arg Tyr Ile His Val
2305 2310 2315 2320
Glu Leu Ile Gly Tyr Pro Pro Pro Ser Ser Ser Ser His Ile Lys Ile
2325 2330 2335
Gly Asp Lys Val Arg Val Lys Ala Ser Val Thr Thr Pro Lys Tyr Lys
2340 2345 2350
Trp Gly Ser Val Thr His Gln Ser Val Gly Val Val Lys Ala Phe Ser
2355 2360 2365
Ala Asn Gly Lys Asp Ile Ile Val Asp Phe Pro Gln Gln Ser His Trp
2370 2375 2380
Thr Gly Leu Leu Ser Glu Met Glu Leu Val Pro Ser Ile His Pro Gly
2385 2390 2395 2400
Val Thr Cys Asp Gly Cys Gln Met Phe Pro Ile Asn Gly Ser Arg Phe
2405 2410 2415
Lys Cys Arg Asn Cys Asp Asp Phe Asp Phe Cys Glu Thr Cys Phe Lys
2420 2425 2430
Thr Lys Lys His Asn Thr Arg His Thr Phe Gly Arg Ile Asn Glu Pro
2435 2440 2445
Gly Gln Ser Ala Val Phe Cys Gly Arg Ser Gly Lys Gln Leu Lys Arg
2450 2455 2460
Cys His Ser Ser Gln Pro Gly Met Leu Leu Asp Ser Trp Ser Arg Met
2465 2470 2475 2480
Val Lys Ser Leu Asn Val Ser Ser Ser Val Asn Gln Ala Ser Arg Leu
2485 2490 2495
Ile Asp Gly Ser Glu Pro Cys Trp Gln Ser Ser Gly Ser Gln Gly Lys
2500 2505 2510
His Trp Ile Arg Leu Glu Ile Phe Pro Asp Val Leu Val His Arg Leu
2515 2520 2525
Lys Met Ile Val Asp Pro Ala Asp Ser Ser Tyr Met Pro Ser Leu Val
2530 2535 2540
Val Val Ser Gly Gly Asn Ser Leu Asn Asn Leu Ile Glu Leu Lys Thr
2545 2550 2555 2560
Ile Asn Ile Asn Pro Ser Asp Thr Thr Val Pro Leu Leu Asn Asp Tyr
2565 2570 2575
Thr Glu Tyr His Arg Tyr Ile Glu Ile Ala Ile Lys Gln Cys Arg Ser
2580 2585 2590
Ser Gly Ile Asp Cys Lys Ile His Gly Leu Ile Leu Leu Gly Arg Ile
2595 2600 2605
Arg Ala Glu Glu Glu Asp Leu Ala Ala Val Pro Phe Leu Ala Ser Asp
2610 2615 2620
Asn Glu Glu Glu Glu Asp Glu Lys Gly Asn Ser Gly Ser Leu Ile Arg
2625 2630 2635 2640
Lys Lys Ala Ala Gly Leu Glu Ser Ala Ala Thr Ile Arg Thr Lys Val
2645 2650 2655
Phe Val Trp Gly Leu Asn Asp Lys Asp Gln Leu Gly Gly Leu Lys Gly
2660 2665 2670
Ser Lys Ile Lys Val Pro Ser Phe Ser Glu Thr Leu Ser Ala Leu Asn
2675 2680 2685
Val Val Gln Val Ala Gly Gly Ser Lys Ser Leu Phe Ala Val Thr Val

2690 2695 2700
Glu Gly Lys Val Tyr Ala Cys Gly Glu Ala Thr Asn Gly Arg Leu Gly
2705 2710 2715 2720
Leu Gly Ile Ser Ser Gly Thr Val Pro Ile Pro Arg Gln Ile Thr Ala
2725 2730 2735
Leu Ser Ser Tyr Val Val Lys Lys Val Ala Val His Ser Gly Gly Arg
2740 2745 2750
His Ala Thr Ala Leu Thr Val Asp Gly Lys Val Phe Ser Trp Gly Glu
2755 2760 2765
Gly Asp Asp Gly Lys Leu Gly His Phe Ser Arg Met Asn Cys Asp Lys
2770 2775 2780
Pro Arg Leu Ile Glu Ala Leu Lys Thr Lys Arg Ile Arg Asp Ile Ala
2785 2790 2795 2800
Cys Gly Ser Ser His Ser Ala Ala Leu Thr Ser Ser Gly Glu Leu Tyr
2805 2810 2815
Thr Trp Gly Leu Gly Glu Tyr Gly Arg Leu Gly His Gly Asp Asn Thr
2820 2825 2830
Thr Gln Leu Lys Pro Lys Met Val Lys Val Leu Leu Gly His Arg Val
2835 2840 2845
Ile Gln Val Ala Cys Gly Ser Arg Asp Ala Gln Thr Leu Ala Leu Thr
2850 2855 2860
Asp Glu Gly Leu Val Phe Ser Trp Gly Asp Gly Asp Phe Gly Lys Leu
2865 2870 2875 2880
Gly Arg Gly Gly Ser Glu Gly Cys Asn Ile Pro Gln Asn Ile Glu Arg
2885 2890 2895
Leu Asn Gly Gln Gly Val Cys Gln Ile Glu Cys Gly Ala Gln Phe Ser
2900 2905 2910
Leu Ala Leu Thr Lys Ser Gly Val Val Trp Thr Trp Gly Lys Gly Asp
2915 2920 2925
Tyr Phe Arg Leu Gly His Gly Ser Asp Val His Val Arg Lys Pro Gln
2930 2935 2940
Val Val Glu Gly Leu Arg Gly Lys Lys Ile Val His Val Ala Val Gly
2945 2950 2955 2960
Ala Leu His Cys Leu Ala Val Thr Asp Ser Gly Gln Val Tyr Ala Trp
2965 2970 2975
Gly Asp Asn Asp His Gly Gln Gln Gly Asn Gly Thr Thr Thr Val Asn
2980 2985 2990
Arg Lys Pro Thr Leu Val Gln Gly Leu Glu Gly Gln Lys Ile Thr Arg
2995 3000 3005
Val Ala Cys Gly Ser Ser His Ser Val Ala Trp Thr Thr Val Asp Val
3010 3015 3020
Ala Thr Pro Ser Val His Glu Pro Val Leu Phe Gln Thr Ala Arg Asp
3025 3030 3035 3040
Pro Leu Gly Ala Ser Tyr Leu Gly Val Pro Ser Asp Ala Asp Ser Ser
3045 3050 3055
Ala Ala Ser Asn Lys Ile Ser Gly Ala Ser Asn Ser Lys Pro Asn Arg
3060 3065 3070
Pro Ser Leu Ala Lys Ile Leu Leu Ser Leu Asp Gly Asn Leu Ala Lys
3075 3080 3085
Gln Gln Ala Leu Ser His Ile Leu Thr Ala Leu Gln Ile Met Tyr Ala
3090 3095 3100
Arg Asp Ala Val Val Gly Ala Leu Met Pro Ala Ala Met Ile Ala Pro
3105 3110 3115 3120
Val Glu Cys Pro Ser Phe Ser Ser Ala Ala Pro Ser Asp Ala Ser Ala
3125 3130 3135
Met Ala Ser Pro Met Asn Gly Glu Glu Cys Met Leu Ala Val Asp Ile
3140 3145 3150
Glu Asp Arg Leu Ser Pro Asn Pro Trp Gln Glu Lys Arg Glu Ile Val
3155 3160 3165

Ser Ser Glu Asp Ala Val Thr Pro Ser Ala Val Thr Pro Ser Ala Pro
3170 3175 3180
Ser Ala Ser Ala Arg Pro Phe Ile Pro Val Thr Asp Asp Leu Gly Ala
3185 3190 3195 3200
Ala Ser Ile Ile Ala Glu Thr Met Thr Lys Thr Lys Glu Asp Val Glu
3205 3210 3215
Ser Gln Asn Lys Ala Ala Gly Pro Glu Pro Gln Ala Leu Asp Glu Phe
3220 3225 3230
Thr Ser Leu Leu Ile Ala Asp Asp Thr Arg Val Val Val Asp Leu Leu
3235 3240 3245
Lys Leu Ser Val Cys Ser Arg Ala Gly Asp Arg Gly Arg Asp Val Leu
3250 3255 3260
Ser Ala Val Leu Ser Gly Met Gly Thr Ala Tyr Pro Gln Val Ala Asp
3265 3270 3275 3280
Met Leu Leu Glu Leu Cys Val Thr Glu Leu Glu Asp Val Ala Thr Asp
3285 3290 3295
Ser Gln Ser Gly Arg Leu Ser Ser Gln Pro Val Val Val Glu Ser Ser
3300 3305 3310
His Pro Tyr Thr Asp Asp Thr Ser Thr Ser Gly Thr Val Lys Ile Pro
3315 3320 3325
Gly Ala Glu Gly Leu Arg Val Glu Phe Asp Arg Gln Cys Ser Thr Glu
3330 3335 3340
Arg Arg His Asp Pro Leu Thr Val Met Asp Gly Val Asn Arg Ile Val
3345 3350 3355 3360
Ser Val Arg Ser Gly Arg Glu Trp Ser Asp Trp Ser Ser Glu Leu Arg
3365 3370 3375
Ile Pro Gly Asp Glu Leu Lys Trp Lys Phe Ile Ser Asp Gly Ser Val
3380 3385 3390
Asn Gly Trp Gly Trp Arg Phe Thr Val Tyr Pro Ile Met Pro Ala Ala
3395 3400 3405
Gly Pro Lys Glu Leu Leu Ser Asp Arg Cys Val Leu Ser Cys Pro Ser
3410 3415 3420
Met Asp Leu Val Thr Cys Leu Leu Asp Phe Arg Leu Asn Leu Ala Ser
3425 3430 3435 3440
Asn Arg Ser Ile Val Pro Arg Leu Ala Ala Ser Leu Ala Ala Cys Ala
3445 3450 3455
Gln Leu Ser Ala Leu Ala Ala Ser His Arg Met Trp Ala Leu Gln Arg
3460 3465 3470
Leu Arg Lys Leu Leu Thr Thr Glu Phe Gly Gln Ser Ile Asn Ile Asn
3475 3480 3485
Arg Leu Leu Gly Glu Asn Asp Gly Glu Thr Arg Ala Leu Ser Phe Thr
3490 3495 3500
Gly Ser Ala Leu Ala Ala Leu Val Lys Gly Leu Pro Glu Ala Leu Gln
3505 3510 3515 3520
Arg Gln Phe Glu Tyr Glu Asp Pro Ile Val Arg Gly Gly Lys Gln Leu
3525 3530 3535
Leu His Ser Pro Phe Phe Lys Val Leu Val Ala Leu Ala Cys Asp Leu
3540 3545 3550
Glu Leu Asp Thr Leu Pro Cys Cys Ala Glu Thr His Lys Trp Ala Trp
3555 3560 3565
Phe Arg Arg Tyr Cys Met Ala Ser Arg Val Ala Val Ala Leu Asp Lys
3570 3575 3580
Arg Thr Pro Leu Pro Arg Leu Phe Leu Asp Glu Val Ala Lys Lys Ile
3585 3590 3595 3600
Arg Glu Leu Met Ala Asp Ser Glu Asn Met Asp Val Leu His Glu Ser
3605 3610 3615
His Asp Ile Phe Lys Arg Glu Gln Asp Glu Gln Leu Val Gln Trp Met
3620 3625 3630
Asn Arg Arg Pro Asp Asp Trp Thr Leu Ser Ala Gly Gly Ser Gly Thr

```
        3635                3640                3645
Ile Tyr Gly Trp Gly His Asn His Arg Gly Gln Leu Gly Gly Ile Glu
    3650                3655                3660
Gly Ala Lys Val Lys Val Pro Thr Pro Cys Glu Ala Leu Ala Thr Leu
3665                3670                3675                3680
Arg Pro Val Gln Leu Ile Gly Gly Glu Gln Thr Leu Phe Ala Val Thr
                3685                3690                3695
Ala Asp Gly Lys Leu Tyr Ala Thr Gly Tyr Gly Ala Gly Gly Arg Leu
            3700                3705                3710
Gly Ile Gly Gly Thr Glu Ser Val Ser Thr Pro Thr Leu Leu Glu Ser
        3715                3720                3725
Ile Gln His Val Phe Ile Lys Lys Val Ala Val Asn Ser Gly Gly Lys
    3730                3735                3740
His Cys Leu Ala Leu Ser Ser Glu Gly Glu Val Tyr Ser Trp Gly Glu
3745                3750                3755                3760
Ala Glu Asp Gly Lys Leu Gly His Gly Asn Arg Ser Pro Cys Asp Arg
                3765                3770                3775
Pro Arg Val Ile Glu Ser Leu Arg Gly Ile Glu Val Val Asp Val Ala
            3780                3785                3790
Ala Gly Gly Ala His Ser Ala Cys Val Thr Ala Ala Gly Asp Leu Tyr
        3795                3800                3805
Thr Trp Gly Lys Gly Arg Tyr Gly Arg Leu Gly His Ser Asp Ser Glu
    3810                3815                3820
Asp Gln Leu Lys Pro Lys Leu Val Glu Ala Leu Gln Gly His Arg Val
3825                3830                3835                3840
Val Asp Ile Ala Cys Gly Ser Gly Asp Ala Gln Thr Leu Cys Leu Thr
                3845                3850                3855
Asp Asp Asp Thr Val Trp Ser Trp Gly Asp Gly Asp Tyr Gly Lys Leu
            3860                3865                3870
Gly Arg Gly Gly Ser Asp Gly Cys Lys Val Pro Met Lys Ile Asp Ser
        3875                3880                3885
Leu Thr Gly Leu Gly Val Val Lys Val Glu Cys Gly Ser Gln Phe Ser
    3890                3895                3900
Val Ala Leu Thr Lys Ser Gly Ala Val Tyr Thr Trp Gly Lys Gly Asp
3905                3910                3915                3920
Tyr His Arg Leu Gly His Gly Ser Asp Asp His Val Arg Arg Pro Arg
                3925                3930                3935
Gln Val Gln Gly Leu Gln Gly Lys Lys Val Ile Ala Ile Ala Thr Gly
            3940                3945                3950
Ser Leu His Cys Val Cys Cys Thr Glu Asp Gly Glu Val Tyr Thr Trp
        3955                3960                3965
Gly Asp Asn Asp Glu Gly Gln Leu Gly Asp Gly Thr Thr Asn Ala Ile
    3970                3975                3980
Gln Arg Pro Arg Leu Val Ala Ala Leu Gln Gly Lys Lys Val Asn Arg
3985                3990                3995                4000
Val Ala Cys Gly Ser Ala His Thr Leu Ala Trp Ser Thr Ser Lys Pro
                4005                4010                4015
Ala Ser Ala Gly Lys Leu Pro Ala Gln Val Pro Met Glu Tyr Asn His
            4020                4025                4030
Leu Gln Glu Ile Pro Ile Ile Ala Leu Arg Asn Arg Leu Leu Leu Leu
        4035                4040                4045
His His Leu Ser Glu Leu Phe Cys Pro Cys Ile Pro Met Phe Asp Leu
    4050                4055                4060
Glu Gly Ser Leu Asp Glu Thr Gly Leu Gly Pro Ser Val Gly Phe Asp
4065                4070                4075                4080
Thr Leu Arg Gly Ile Leu Ile Ser Gln Gly Lys Glu Ala Ala Phe Arg
                4085                4090                4095
Lys Val Val Gln Ala Thr Met Val Arg Asp Arg Gln His Gly Pro Val
            4100                4105                4110
```

```
Val Glu Leu Asn Arg Ile Gln Val Lys Arg Ser Arg Ser Lys Gly Gly
        4115                4120                4125
Leu Ala Gly Pro Asp Gly Thr Lys Ser Val Phe Gly Gln Met Cys Ala
    4130                4135                4140
Lys Met Ser Ser Phe Gly Pro Asp Ser Leu Leu Leu Pro His Arg Val
4145                4150                4155                4160
Trp Lys Val Lys Phe Val Gly Glu Ser Val Asp Asp Cys Gly Gly Gly
                4165                4170                4175
Tyr Ser Glu Ser Ile Ala Glu Ile Cys Glu Glu Leu Gln Asn Gly Leu
            4180                4185                4190
Thr Pro Leu Leu Ile Val Thr Pro Asn Gly Arg Asp Glu Ser Gly Ala
        4195                4200                4205
Asn Arg Asp Cys Tyr Leu Leu Ser Pro Ala Ala Arg Ala Pro Val His
    4210                4215                4220
Ser Ser Met Phe Arg Phe Leu Gly Val Leu Leu Gly Ile Ala Ile Arg
4225                4230                4235                4240
Thr Gly Ser Pro Leu Ser Leu Asn Pro Cys Arg Ala Leu Ser Gly Ser
                4245                4250                4255
Ser Trp Leu Gly  *
            4260
```

```
<210> 1356
<211> 64
<212> PRT
<213> Homo sapiens

<400> 1356
Met Ser Lys Val Lys Pro Leu His Gly Ala Pro Ala Pro Leu Leu Val
 1               5                  10                  15
Ser Leu Cys Leu Leu Ser Trp Cys Gly Leu Pro Gly Val Ile Val His
            20                  25                  30
Val Thr Tyr Val Ser Pro Arg His Leu Ser Asn Thr Arg Ser Gly Leu
        35                  40                  45
Glu Ser Ile His Gly Cys Asp Pro Met His Gly Ser Pro Val Gly  *
    50                  55                  60          63
```

```
<210> 1357
<211> 111
<212> PRT
<213> Homo sapiens

<221> misc_feature
<222> (1)...(111)
<223> Xaa = any amino acid or nothing

<400> 1357
Met Ile Phe Asn Lys Ala Ala Asp Thr Leu Gly Asp Val Trp Ile Leu
 1               5                  10                  15
Leu Ala Thr Leu Lys Val Leu Ser Leu Leu Trp Leu Leu Tyr Tyr Val
            20                  25                  30
Ala Ser Thr Thr Arg Gln Pro His Ala Val Leu Tyr Gln Asp Pro His
        35                  40                  45
Ala Gly Pro Leu Trp Val Arg Ser Ser Leu Val Leu Phe Gly Ser Cys
```

```
    50                  55                  60
Thr Phe Cys Leu Asn Ile Phe Arg Val Gly Tyr Asp Val Ser His Ile
65                  70                  75                  80
Arg Cys Lys Ser Gln Leu Asp Leu Val Phe Pro Val Ile Glu Met Val
                85                  90                  95
Phe Ile Gly Val Gln Thr Cys Val Leu Trp Lys His Cys Arg Xaa
            100                 105                 110 111
```

```
<210> 1358
<211> 47
<212> PRT
<213> Homo sapiens
```

```
<400> 1358
Met Ala Leu Leu Ile Ser Thr Cys Ile Asn Lys Ala Val Leu Arg Phe
 1               5                  10                  15
Thr Leu Ser Ser Met Asn Asn Lys Ile Ile Leu Ser Trp Tyr Ser Phe
            20                  25                  30
Asn Val Ile Leu Ile Phe His Glu Asn Val Val Tyr Tyr Ile  *
        35                  40                  45  46
```

```
<210> 1359
<211> 73
<212> PRT
<213> Homo sapiens
```

```
<400> 1359
Met Phe Ser Pro Cys Gly Pro Ala Ser Leu Gly Leu Leu Phe Val Leu
 1               5                  10                  15
Cys Thr His Ser Gln Ala Leu Ala Phe Phe Trp Gly Pro Ser Ser Leu
            20                  25                  30
Ile Gly Ala Ser Gly Phe Leu Leu Gln Arg Thr Ser Leu Leu Arg His
        35                  40                  45
Val Phe Leu Gly Leu Val Tyr Ala Cys Trp Ala His Trp Leu Tyr Cys
    50                  55                  60
Ser Ser Arg Pro Val Thr Lys Glu  *
65                  70      72
```

```
<210> 1360
<211> 57
<212> PRT
<213> Homo sapiens
```

```
<400> 1360
Met Lys Thr Gly Ser Leu Leu Leu Thr Leu Trp Phe Ser Gln Thr Phe
 1               5                  10                  15
Ser Phe Asn Leu Phe Phe Ala Pro Pro His Ser Leu Leu Gln Ser Ser
            20                  25                  30
Ile Phe Phe Ser Val Ser Ser Ile Thr Thr Val His Pro Ile Leu Val
        35                  40                  45
```

Phe Phe Phe Ala Phe Phe Arg Thr *
50 55 56

<210> 1361
<211> 77
<212> PRT
<213> Homo sapiens

<400> 1361
Met Phe Val Leu Phe Leu Ile Leu Val Leu Arg Asn His Phe Leu Val
1 5 10 15
Thr Ile Lys Tyr Gly Val Gly Cys Gly Phe Ile Ile Ser Val Cys Leu
20 25 30
Arg Ala Lys His Phe Asn Phe Asp Glu Ala Gln Phe Val Ser Phe Phe
35 40 45
Leu Cys Asp Ser Cys Phe Cys Leu Leu Arg Asn Leu Pro Thr Gln Arg
50 55 60
Leu Gln Arg Phe Phe Phe Cys Trp Phe Phe Leu Ile *
65 70 75 76

<210> 1362
<211> 106
<212> PRT
<213> Homo sapiens

<400> 1362
Met Gln Asn Arg Thr Gly Leu Ile Leu Cys Ala Leu Ala Leu Leu Met
1 5 10 15
Gly Phe Leu Met Val Cys Leu Gly Ala Phe Phe Ile Ser Trp Gly Ser
20 25 30
Ile Phe Asp Cys Gln Gly Ser Leu Ile Ala Ala Tyr Leu Leu Leu Pro
35 40 45
Leu Gly Phe Val Ile Leu Leu Ser Gly Ile Phe Trp Ser Asn Tyr Arg
50 55 60
Gln Val Thr Glu Ser Lys Gly Val Leu Arg His Met Leu Arg Gln His
65 70 75 80
Leu Ala His Gly Ala Leu Pro Val Ala Thr Val Asp Arg Ala Ala Leu
85 90 95
Leu Lys Ile Met Cys Lys Gln Leu Leu *
100 105

<210> 1363
<211> 57
<212> PRT
<213> Homo sapiens

<400> 1363
Met Ala Trp Lys Pro Leu Gly Arg Gln Ala Val Leu Arg Glu Thr Pro
1 5 10 15
Leu Ala Thr Leu Cys Ile Asp Arg Arg Gln Val Ser Ser Ser Leu Val

```
                20                  25                  30
Gln Glu Gly Phe His Ser Lys Ser Cys His Cys Leu Gly Asp Ser Phe
            35                  40                  45
Arg Glu Lys Asn Gln Val Val Gly  *
    50                  55  56


     <210> 1364
     <211> 75
     <212> PRT
     <213> Homo sapiens


     <400> 1364
Met Cys Leu Leu Lys Ala Ala Pro Phe Phe Phe Phe Tyr Val Pro Gln
 1               5                  10                  15
Val Gly Lys Gly Asn Pro Arg Pro Pro Arg Gly Cys Ser Ala Phe His
                20                  25                  30
Pro Pro Thr His Leu Arg Pro Gly Ser Cys Ser Val Ala Gln Ala Gly
            35                  40                  45
Val Gln Trp Arg Ser Leu Gly Ser Ile Ala Ala Ser Val Ser Trp Val
    50                  55                  60
Gln Ala Ile Leu Leu Pro Gln Pro Leu Glu  *
65                  70              74


     <210> 1365
     <211> 58
     <212> PRT
     <213> Homo sapiens


     <400> 1365
Met Lys Leu Gln Val Phe Ala Val Asn Ile Thr Ala Leu Lys Ala Ala
 1               5                  10                  15
Arg Leu Glu Leu Phe Val Leu Pro Gly Gly Phe Ile Val Phe Leu Ala
                20                  25                  30
Ser Glu Leu Lys Leu Gln Thr Ser Leu Glu Ser Val Ala Pro His Lys
            35                  40                  45
Asp Ser Met Ser Leu Lys Ser Glu His  *
    50                  55      57


     <210> 1366
     <211> 58
     <212> PRT
     <213> Homo sapiens


     <400> 1366
Met His Cys Ser Phe Ile Ser Ala Phe Leu Leu Pro Val Phe Leu Ser
 1               5                  10                  15
Leu Thr Val Ser Ala Ser Ile Phe Val Ser Leu His Ser Phe Pro Leu
                20                  25                  30
Ser Leu Ser Tyr Phe Ser Phe Leu Gly Ser Phe Phe Leu Ser Val Cys
            35                  40                  45
```

```
Leu Asp Leu Tyr Ser Ser Leu Phe Phe  *
     50                 55      57


     &lt;210&gt; 1367
     &lt;211&gt; 48
     &lt;212&gt; PRT
     &lt;213&gt; Homo sapiens


     &lt;400&gt; 1367
Met Met Gly Arg Ile Phe Ala Ala Leu Ser Leu Ile Lys Leu Met Met
 1               5                  10                  15
Tyr Ser Leu Phe Pro Val Ile Glu Ser Ser Leu Cys His Leu Glu Val
            20                  25                  30
Trp Ala Trp Arg His Ile Trp Pro Thr Ala Gly Arg Gly Val Pro  *
        35                  40                  45      47


     &lt;210&gt; 1368
     &lt;211&gt; 96
     &lt;212&gt; PRT
     &lt;213&gt; Homo sapiens


     &lt;400&gt; 1368
Met Gly Arg Arg Lys Ser Phe Phe Phe Leu Phe Leu Glu Cys Arg Gln
 1               5                  10                  15
Lys Gly Leu His Ile Pro Leu Cys Thr Cys Ser His Ala Pro Arg Pro
            20                  25                  30
Pro Leu Ala Ala Pro Ser Ala Leu Ile Leu Pro Pro Glu Ile Ser His
        35                  40                  45
Thr Ser Arg Gly Ile Leu Leu Ser His Gly Leu Phe Pro Thr Ala Thr
    50                  55                  60
Met Pro Leu Phe Phe Pro Ser His Ala Ser His Ser Pro Thr Val Thr
65                  70                  75                  80
Met Pro Leu Phe Phe Pro Ser His Ala Ser His Ser Pro Ser Thr  *
                85                  90                  95


     &lt;210&gt; 1369
     &lt;211&gt; 76
     &lt;212&gt; PRT
     &lt;213&gt; Homo sapiens


     &lt;400&gt; 1369
Met Trp Asp His Phe Ile Leu Ser Arg Val Leu Phe Cys Leu Phe Val
 1               5                  10                  15
Phe His Ser Arg Val Leu Lys Asp His Met Ala Ser Asn Ala Tyr Lys
            20                  25                  30
Ser Ala Leu Phe Phe Thr Val Arg Tyr Leu Glu Thr Lys Gln Phe Leu
        35                  40                  45
Leu Arg Cys Cys Cys Trp Pro Asp Ala Val Ala His Ala Cys Asn Thr
    50                  55                  60
Ser Thr Leu Arg Gly Gln Gly Arg His Ile Thr  *
```

65 70 75

<210> 1370
<211> 79
<212> PRT
<213> Homo sapiens

<400> 1370
Met Cys Ser Cys Leu His Thr Leu Gln Arg Arg Phe Leu His Phe Val
1 5 10 15
Ser Ile Ala Leu Ser Lys Ile Trp Gln Asn Asn Ala Phe His Leu Gln
20 25 30
Val Glu Val Ser Trp Leu Ser Thr Phe Val Asp Lys Val Ile Val Met
35 40 45
Arg Leu Ile Ser Ser Lys His Phe Thr Asp Thr Met Asn Asp Arg Val
50 55 60
His Ser Phe Leu Asn Asp Ile Gly Phe Val Cys Leu Leu Ser *
65 70 75 78

<210> 1371
<211> 227
<212> PRT
<213> Homo sapiens

<221> misc_feature
<222> (1)...(227)
<223> Xaa = any amino acid or nothing

<400> 1371
Met Leu Tyr Phe Gln Leu Val Ile Met Ala Gly Thr Val Leu Leu Ala
1 5 10 15
Tyr Tyr Phe Glu Cys Thr Asp Thr Phe Gln Val His Ile Gln Gly Phe
20 25 30
Phe Cys Gln Asp Gly Asp Leu Met Lys Pro Tyr Pro Gly Thr Glu Glu
35 40 45
Glu Ser Phe Ile Thr Pro Leu Val Leu Tyr Cys Val Leu Ala Ala Thr
50 55 60
Pro Thr Ala Ile Ile Phe Ile Gly Glu Ile Ser Met Tyr Phe Ile Lys
65 70 75 80
Ser Thr Arg Glu Ser Leu Ile Ala Gln Glu Lys Thr Ile Leu Thr Gly
85 90 95
Glu Cys Cys Tyr Leu Asn Pro Leu Leu Arg Arg Ile Ile Arg Phe Thr
100 105 110
Gly Val Phe Ala Phe Gly Leu Phe Ala Thr Asp Ile Phe Val Asn Ala
115 120 125
Gly Gln Val Val Thr Gly His Leu Thr Pro Tyr Phe Leu Thr Val Cys
130 135 140
Lys Pro Asn Tyr Thr Ser Ala Asp Cys Gln Ala His His Gln Phe Ile
145 150 155 160
Asn Asn Gly Asn Ile Cys Thr Gly Asp Leu Gly Ser Asp Arg Lys Gly
165 170 175
Ser Glu Ile Leu Ser Leu Gln Thr Arg Cys Ser Glu His Leu Leu Arg
180 185 190

```
Leu Ile Trp Pro Arg Cys Ile Phe Thr Arg His Asn Gln Gly Arg Gly
        195                 200                 205
Gly Ser Ser Met Gly Pro Ser Arg Trp Leu Cys Leu Gly Thr Phe Leu
    210                 215                 220
His Xaa Leu
225     227
```

<210> 1372
<211> 99
<212> PRT
<213> Homo sapiens

<400> 1372

```
Met Phe Leu Ser Leu Ser Leu Thr Leu Cys Leu Cys Phe Ser Phe Phe
  1               5                  10                  15
Cys Leu Tyr Leu Ser Leu Ser Leu Tyr Leu Arg Ser Phe Phe Cys Leu
             20                  25                  30
Pro Phe His Val Ser Val Phe Leu Cys Leu Phe Pro Ser Val Leu Phe
         35                  40                  45
Leu Ser Val Ala Leu Gly Ser Pro Glu Asn His Ile Ser Trp Arg Lys
     50                  55                  60
Val Gly Glu Glu Leu Lys Leu Ala Ser His Arg Asn Phe Cys Ser Leu
 65                  70                  75                  80
Ile Gln Met Met Arg Ser Asn Lys Pro Ser Pro Ser Arg Gln Arg Gly
                 85                  90                  95
Trp Ala  *
     98
```

<210> 1373
<211> 69
<212> PRT
<213> Homo sapiens

<400> 1373

```
Met Leu His Thr Pro Gln Thr Cys Arg Pro Gly Leu Cys Val Leu Ala
  1               5                  10                  15
Ser Arg Pro Val Leu Tyr Thr Leu Cys Leu Leu Ile Pro Val Leu Cys
             20                  25                  30
Gly Asp Thr Phe Trp Ala Ser Trp Ser Leu Leu Thr Lys Ala Thr Pro
         35                  40                  45
Ser Ser Leu Leu Cys Leu Ser Asp Lys Ser Ile Pro Ser Leu Ile Ser
     50                  55                  60
Lys Gly Asp Ser  *
 65          68
```

<210> 1374
<211> 296
<212> PRT
<213> Homo sapiens

<400> 1374
Met Arg Ser Lys Ile Met Ile His Ile His Ile Phe Leu Leu Ala Ser
1 5 10 15
Phe Arg Phe Lys Glu His Val Gln Asn Asn Leu Pro Arg Asp Leu Leu
20 25 30
Thr Gly Glu Gln Phe Ile Gln Leu Arg Arg Glu Leu Ala Ser Val Asn
35 40 45
Gly His Ser Gly Asp Asp Gly Pro Pro Gly Asp Asp Leu Pro Ser Gly
50 55 60
Ile Glu Asp Ile Thr Asp Pro Ala Lys Leu Ile Thr Glu Ile Glu Asn
65 70 75 80
Met Arg His Arg Ile Ile Glu Ile His Gln Glu Met Phe Asn Tyr Asn
85 90 95
Glu His Glu Val Ser Lys Arg Trp Thr Phe Glu Glu Gly Ile Lys Arg
100 105 110
Pro Tyr Phe His Val Lys Pro Leu Glu Lys Ala Gln Leu Lys Asn Trp
115 120 125
Lys Glu Tyr Leu Glu Phe Glu Ile Glu Asn Gly Thr His Glu Arg Val
130 135 140
Val Val Leu Phe Glu Arg Cys Val Ile Ser Cys Ala Leu Tyr Glu Glu
145 150 155 160
Phe Trp Ile Lys Tyr Ala Lys Tyr Met Glu Asn His Ser Ile Glu Gly
165 170 175
Val Arg His Val Phe Ser Arg Ala Cys Thr Ile His Leu Pro Lys Lys
180 185 190
Pro Met Val His Met Leu Trp Ala Ala Phe Glu Glu Gln Gln Gly Asn
195 200 205
Ile Asn Glu Ala Arg Asn Ile Leu Lys Thr Phe Glu Glu Cys Val Leu
210 215 220
Gly Leu Ala Met Val Arg Leu Arg Arg Val Ser Leu Glu Arg Arg His
225 230 235 240
Gly Asn Leu Glu Glu Ala Glu His Leu Leu Gln Asp Ala Ile Lys Asn
245 250 255
Ala Lys Ser Asn Asn Glu Ser Ser Phe Tyr Ala Val Lys Leu Ala Arg
260 265 270
His Leu Phe Lys Ile Gln Lys Asn Leu Pro Lys Ser Arg Lys Val Leu
275 280 285
Leu Glu Ala Ile Glu Arg Asp Lys
290 295 296

<210> 1375
<211> 75
<212> PRT
<213> Homo sapiens

<400> 1375
Met Cys Leu Leu Lys Ala Ala Pro Phe Phe Phe Phe Tyr Val Pro Gln
1 5 10 15
Val Gly Lys Gly Asn Pro Arg Pro Pro Arg Gly Cys Ser Ala Phe His
20 25 30
Pro Pro Thr His Leu Arg Pro Gly Ser Cys Ser Val Ala Gln Ala Gly
35 40 45
Val Gln Trp Arg Ser Leu Gly Ser Ile Ala Ala Ser Val Ser Trp Val
50 55 60
Gln Ala Ile Leu Leu Pro Gln Pro Leu Glu *
65 70 74

<210> 1376
<211> 61
<212> PRT
<213> Homo sapiens

<400> 1376
```
Met Cys Tyr Glu Trp Val Ile Thr Thr Val Gly Ser Trp Ala Leu Leu
 1               5                  10                  15
Cys Gln Arg Thr Leu Trp Lys Pro His Arg Thr Tyr Gln Lys Leu Thr
            20                  25                  30
Leu Asn Ser Cys Pro Thr Pro Ile Val Glu Gly Gly Leu Glu Ser Phe
        35                  40                  45
Pro Ser Pro Asn Phe Pro Ser Cys Ile Ser Trp Ser  *
    50                  55                  60
```

<210> 1377
<211> 110
<212> PRT
<213> Homo sapiens

<400> 1377
```
Met Trp Val Trp Val Thr Ala Ala His Leu Leu Cys Ser Leu Ala Ala
 1               5                  10                  15
Ser Phe Val Lys Lys Lys Ser Leu Gly Lys Leu Arg Val Asp Val Cys
            20                  25                  30
Arg Ser Pro Pro Pro Glu Gly Ser Arg Thr Gln Thr Ser Ser Ser Leu
        35                  40                  45
Phe Tyr Arg Gly Gly Asn Gly Ala Ser Tyr Ala Asn Tyr Ile Leu His
    50                  55                  60
His Thr Met Ala Leu Glu Gly Gln Arg Ser His Trp Ala Pro Cys Val
65                  70                  75                  80
Ser Cys Pro Ala Gln Gly Leu Ala Leu Arg Arg Gly Cys Thr Thr Phe
                85                  90                  95
Leu His Lys Asn Lys Gly Gly Thr Glu Ala Val Thr Val  *
            100                 105                 109
```

<210> 1378
<211> 47
<212> PRT
<213> Homo sapiens

<400> 1378
```
Met Phe Ala Leu Gln Lys Met Arg Leu Cys Val Leu Trp Arg Val Leu
 1               5                  10                  15
Glu Glu Gly Gly Ile Thr Arg Phe Gly Asp Ser His Ser Asp Ser Leu
            20                  25                  30
Leu Phe Ser Val Thr Phe Arg Ile His Arg Asp Met Phe Cys  *
        35                  40                  45  46
```

<210> 1379
<211> 140
<212> PRT
<213> Homo sapiens

<400> 1379
Met Arg His Pro Ser Pro Trp Pro Phe Leu Phe Phe Cys Phe Val Pro
1 5 10 15
Ala Thr Leu Arg Ser Phe Pro Ser Gly Leu Val Trp Pro Gly Cys Trp
20 25 30
Trp Glu Pro Arg Ala Ser Pro Ser Ser Leu Ala Pro Gly Met Lys Ser
35 40 45
Gln Leu Trp Ala Ala Ala Trp Arg Pro Gly Thr Ser Leu Gln Gly Met
50 55 60
Ala Gly Ile Leu Arg Gln Ala Ala Glu Ala Gly Pro Ala Gly Val Ala
65 70 75 80
Leu Ile Leu Ile Lys Gly Thr Gly Asn Glu Glu Pro Leu Gly Pro Leu
85 90 95
Pro Ser Arg Cys Leu Cys Pro Pro Pro Glu Glu Pro Arg Phe His Trp
100 105 110
Ala Leu Gly Lys Glu Pro Thr Gly Pro Gly Arg Pro Gln Pro Val Gln
115 120 125
His His Ile Glu Gly Pro His Pro Val Gly Phe Gly
130 135 140

<210> 1380
<211> 50
<212> PRT
<213> Homo sapiens

<400> 1380
Met Gln Glu Pro Leu Thr Phe Leu Gln Leu Leu Arg Trp Gln Leu Phe
1 5 10 15
Pro Leu Pro Asp Ser Pro Thr Phe Ser Ala Phe Ile Leu Val Gly Leu
20 25 30
Cys Arg Met Leu Phe Ala Gly Arg Ile Ile Ser Gly Leu Thr Arg Val
35 40 45
Ile *
49

<210> 1381
<211> 78
<212> PRT
<213> Homo sapiens

<400> 1381
Met Leu Arg Leu Asp Ile Ile Asn Ser Leu Val Thr Thr Val Phe Met
1 5 10 15
Leu Ile Val Ser Val Leu Ala Leu Ile Pro Glu Thr Thr Thr Leu Thr
20 25 30

Val Gly Gly Gly Val Phe Ala Leu Val Thr Ala Val Cys Cys Leu Ala
35 40 45
Asp Gly Ala Leu Ile Tyr Arg Lys Leu Leu Phe Asn Pro Ser Gly Pro
50 55 60
Tyr Gln Lys Lys Pro Val His Glu Lys Lys Glu Val Leu *
65 70 75 77

<210> 1382
<211> 57
<212> PRT
<213> Homo sapiens

<400> 1382
Met Leu Thr Thr Leu Leu Leu Leu Leu His Lys Arg Ile Phe Arg Gly
1 5 10 15
Asn Phe His Ile Leu His Phe His Ile Cys Ile Gln Ile Lys Lys Gln
20 25 30
Ile Pro Ile Leu Glu Asn Asp Leu Phe Lys Met Tyr Thr Val Ser Asn
35 40 45
Lys Ala Lys Thr Arg Thr Trp Ser *
50 55 56

<210> 1383
<211> 64
<212> PRT
<213> Homo sapiens

<400> 1383
Met Val Cys Arg Leu Pro Cys Thr Leu Leu Pro Trp Pro Leu Lys His
1 5 10 15
Lys Gln Gly Ala Leu Leu Tyr Ile Cys Pro Ala Ser Leu Pro Ala Phe
20 25 30
Asn Pro Arg Asn Leu Ser Val Tyr Leu Leu Phe Ser Ala Ser Glu Ser
35 40 45
Leu Pro Leu Lys Ser Glu Gln Ala Arg Pro Gly Gly Ser Arg Leu *
50 55 60 63

<210> 1384
<211> 67
<212> PRT
<213> Homo sapiens

<400> 1384
Met Leu Ser Phe Val Pro Leu Leu Ser Ser Trp Leu Gly Thr Trp Ile
1 5 10 15
Thr Asp Arg Gly Ala Ala Gly Ser Cys Gln Ala Glu Ala Pro Arg Leu
20 25 30
Ala Gly Glu Thr Ala Gly Gln Arg Val Trp Glu Arg Gly Met Gln Arg
35 40 45
Ala Ala Ala Val Gly Lys Ile Leu Asp Pro Lys Gly His Thr Ala Ser

```
    50                   55                   60
Pro His  *
 65  66


     &lt;210&gt; 1385
     &lt;211&gt; 50
     &lt;212&gt; PRT
     &lt;213&gt; Homo sapiens


     &lt;400&gt; 1385
Met Leu Val Leu Phe Val Ala Thr Trp Ser Asp Leu Gly Leu Cys Lys
 1               5                  10                  15
Lys Arg Pro Lys Pro Gly Gly Trp Asn Thr Gly Gly Cys Arg Tyr Pro
            20                  25                  30
Gly Leu Ala Cys Pro Leu Gly Arg Pro Pro Gly Gln Trp Gly Ala Thr
        35                  40                  45
Val  *
 49


     &lt;210&gt; 1386
     &lt;211&gt; 123
     &lt;212&gt; PRT
     &lt;213&gt; Homo sapiens


     &lt;400&gt; 1386
Met Lys Trp Val Thr Phe Ile Ser Leu Leu Phe Leu Phe Ser Ser Ala
 1               5                  10                  15
Tyr Ser Arg Gly Pro Lys Ala Glu Phe Ala Glu Val Ser Lys Leu Val
            20                  25                  30
Thr Asp Leu Thr Lys Val His Thr Glu Cys Cys His Gly Asp Leu Leu
        35                  40                  45
Glu Cys Ala Asp Asp Arg Ala Asp Leu Ala Lys Tyr Ile Cys Glu Asn
    50                  55                  60
Gln Asp Ser Ile Ser Ser Lys Leu Lys Glu Cys Cys Glu Lys Pro Leu
65                  70                  75                  80
Leu Glu Lys Ser His Cys Ile Ala Glu Val Glu Asn Asp Glu Met Pro
                85                  90                  95
Ala Asp Leu Pro Ser Leu Ala Ala Asp Phe Val Glu Ser Lys Asp Val
            100                 105                 110
Cys Lys Asn Tyr Ala Glu Ala Lys Asp Val Phe
        115                 120         123


     &lt;210&gt; 1387
     &lt;211&gt; 65
     &lt;212&gt; PRT
     &lt;213&gt; Homo sapiens


     &lt;400&gt; 1387
Met Pro Arg Leu Phe Ser Pro Leu Ile Leu Leu His Thr Leu Ser Leu
 1               5                  10                  15
```

Lys Ser His Glu Thr Phe Gln Trp Ser Gln Phe Leu Tyr Gln Asn Thr
20 25 30
Arg Asp Ala Cys Phe Thr Trp Thr Tyr Ile Phe Pro Arg Ile Thr Trp
35 40 45
Ile Asn Glu Trp Cys Cys Phe Pro Val Val Gly Glu Lys Leu Gly Thr
50 55 60 64
*

<210> 1388
<211> 56
<212> PRT
<213> Homo sapiens

<400> 1388
Met Gly Leu Leu Asn Lys Tyr Ala Ser Val Ile Ile Tyr Leu Tyr Phe
1 5 10 15
Ser Leu Val Lys Ser Glu Ser Leu Phe His Leu Met Tyr Leu Pro Ser
20 25 30
Leu Phe Ile Gln Phe Phe Leu Gly Ile Phe Ser Leu Lys Thr His Cys
35 40 45
Cys Thr Ser Lys Phe Asp Ser *
50 55

<210> 1389
<211> 76
<212> PRT
<213> Homo sapiens

<400> 1389
Met Arg Arg Arg Ala Leu Lys His Trp Val Ala Leu Cys Leu Thr Trp
1 5 10 15
Thr Ala Gly Glu Ser Thr Gly Pro Trp Pro Ser Pro Glu Pro Ser Val
20 25 30
Arg Ala Lys Glu Ala Asp Pro Ser Gly Arg Arg Ser Leu Gly Ser Pro
35 40 45
Gly Leu Glu Cys Gly Pro Arg Leu Thr Arg Gly Ser Gly Arg Gln Cys
50 55 60
Asp Gly Pro Arg Gly Ile Cys His Ala Leu Gly *
65 70 75

<210> 1390
<211> 149
<212> PRT
<213> Homo sapiens

<400> 1390
Met Ala Ala Ser Pro Ala Arg Pro Ala Val Leu Ala Leu Thr Gly Leu
1 5 10 15
Ala Leu Leu Leu Leu Leu Cys Trp Gly Pro Gly Gly Ile Ser Gly Asn

```
                    20                  25                  30
Lys Leu Lys Leu Met Leu Gln Lys Arg Glu Ala Pro Val Pro Thr Lys
            35                  40                  45
Thr Lys Val Ala Val Asp Glu Asn Lys Ala Lys Glu Phe Leu Gly Ser
        50                  55                  60
Leu Lys Arg Gln Lys Arg Gln Leu Trp Asp Arg Thr Arg Pro Glu Val
65                  70                  75                  80
Gln Gln Trp Tyr Gln Gln Phe Leu Tyr Met Gly Phe Asp Glu Ala Lys
                85                  90                  95
Phe Glu Asp Asp Ile Thr Tyr Trp Leu Asn Arg Asp Arg Asn Gly His
            100                 105                 110
Glu Tyr Tyr Gly Asp Tyr Tyr Gln Arg His Tyr Asp Glu Asp Ser Ala
        115                 120                 125
Ile Gly Pro Arg Ser Pro Tyr Gly Phe Arg His Gly Ala Ser Val Asn
    130                 135                 140
Tyr Asp Asp Tyr  *
145         148
```

<210> 1391
<211> 125
<212> PRT
<213> Homo sapiens

<400> 1391

```
Met Val Met Gly Trp His Trp Pro Gln Gly Leu Gly Leu Ser Leu Ser
 1               5                  10                  15
Leu Cys Pro Ser Asp Leu Asp Gly Trp Val Ser Arg Glu Val Pro Leu
            20                  25                  30
Leu Asp Arg Pro Gln Ala Leu Pro Pro Cys Val Gln Ile Leu Ser Ala
        35                  40                  45
Pro Ala Ser Thr Ser Cys Pro Ser Ala Leu Ser Pro Trp His Asp Pro
    50                  55                  60
Gly Leu Pro Val Thr Ser Gln Asn His Phe Ala Trp Phe Pro Leu Gly
65                  70                  75                  80
Ser Lys Ala Cys Leu Gly Pro Ser Ile Asp Arg Glu Ala Val Lys Glu
                85                  90                  95
Ile Asn Ala Glu Glu Gly Val Arg Arg Gln Thr Gln Gly Pro Ile Lys
            100                 105                 110
Val Arg Lys Gln Ala Gly Cys Gly Gly Ser Cys Leu  *
        115                 120                 124
```

<210> 1392
<211> 56
<212> PRT
<213> Homo sapiens

<400> 1392

```
Met Ile Ile Gln Ile Cys Thr Ile Ser Arg Ile Glu Phe Ile Cys Leu
 1               5                  10                  15
Cys Val Cys Val Phe Phe Arg Val Ile Trp Leu Pro Val Glu Phe Tyr
            20                  25                  30
Leu Glu Thr Lys Ile Leu Lys Val Val Phe Val Ile Val Phe Val Pro
        35                  40                  45
```

Ile Ile Leu Pro Leu His Pro *
50 55

<210> 1393
<211> 55
<212> PRT
<213> Homo sapiens

<400> 1393
Met Glu Ala Trp Lys Ala Leu Ile Gly Leu Phe Pro Leu Arg Ser Ser
1 5 10 15
Ala Ser Pro Phe Thr Tyr His Cys Trp Glu Pro Ala Gln Pro Ala His
20 25 30
Gln Glu Phe His Ser Thr Ile Ala Leu Arg Gly Arg Gly Gly Lys Pro
35 40 45
Gln Glu Glu Ser Ser Pro *
50 54

<210> 1394
<211> 51
<212> PRT
<213> Homo sapiens

<400> 1394
Met Ser Leu Asn Pro Glu Phe Leu Trp Leu Lys Trp Phe Ser Leu Leu
1 5 10 15
Leu Arg Gly Arg Arg Asn Ser Cys Leu Ile Ala Leu Lys Gly Tyr His
20 25 30
Ser Val Met Ile Phe His Leu Pro Leu Ile Pro Ser Ser Val Thr Ser
35 40 45
Cys His *
50

<210> 1395
<211> 105
<212> PRT
<213> Homo sapiens

<400> 1395
Met Pro Cys Phe Met Pro Asn Pro Gly Ala Val Leu Gly Leu Pro Pro
1 5 10 15
Trp Leu Leu Ser Thr Gln Arg Leu Thr His Thr Arg Ala Tyr Leu Asn
20 25 30
Trp Leu Ala Ser Asp Arg Trp Met Arg Arg His Trp Arg Thr Gly Glu
35 40 45
Ser Gln Val Glu Arg Ser Ser Arg Pro Trp Trp Glu Thr Gln His Leu
50 55 60
Ser Pro Ala Ser Leu Gly Arg Arg Pro Ala Pro Gly Leu Gln Glu His
65 70 75 80
Phe Leu Asp Thr Asp Gly Lys Val Ala Asp Ser Gly Leu Gln Met Gly

```
                    85                  90                  95
Phe Gly Leu Leu Ser Leu Pro Ser Ile
            100                 105


     <210> 1396
     <211> 49
     <212> PRT
     <213> Homo sapiens


     <400> 1396
Met Leu Cys Asn Leu Ala Leu Lys Leu Leu Asn Cys Val Ser Ala Trp
 1               5                  10                  15
Asn Met Asn Ile Arg Leu Lys Cys Leu Leu Lys Pro Lys Asn Val Ser
            20                  25                  30
Lys Val Cys Ser Arg Gly Leu Tyr Phe Ile Tyr Val Met Asp Ser Leu
        35                  40                  45          48
 *


     <210> 1397
     <211> 104
     <212> PRT
     <213> Homo sapiens


     <400> 1397
Met Leu Ser Trp Val Phe Pro Gly Ser Val Phe Gly Leu Cys Leu Ser
 1               5                  10                  15
Val Trp Val Phe Trp His Gln Ala Ser Leu Gly Arg Ala Ser Gly Cys
            20                  25                  30
Ala Pro Ala Leu Arg Val Gly Leu Ile Pro Gly Cys Arg Gly Leu Arg
        35                  40                  45
Ala Glu Leu Phe His Leu Glu Asp Lys Asp Gly Ser Ser Gly Leu Gly
    50                  55                  60
Gly Gly Gly Gly Ala Gly His Asp Leu Ile Leu Arg Arg Ala Trp Cys
65                  70                  75                  80
Trp Gly Leu Thr Asp Asp Gly Glu Ala Arg Val Gln Ala Leu Gly Met
                85                  90                  95
Thr Pro Gly Ile Ala Phe Ser  *
            100         103


     <210> 1398
     <211> 82
     <212> PRT
     <213> Homo sapiens


     <400> 1398
Met Lys Pro Val Trp Val Ala Thr Leu Leu Trp Met Leu Leu Leu Val
 1               5                  10                  15
Pro Arg Leu Gly Ala Ala Arg Lys Gly Ser Pro Glu Glu Ala Ser Phe
            20                  25                  30
```

```
Tyr Tyr Gly Thr Phe Pro Leu Gly Gly His His Ser Ala Glu Gly Thr
        35                  40                  45
Ala Arg Gln Pro Leu Pro Ile Leu Pro Val Leu Ala Pro Ala Pro Ala
    50                  55                  60
His Arg His Pro Ser Arg Ala Gly Glu Gln Glu Gly Asn Arg Ile Leu
65                  70                  75                  80
Gln  *
81
```

<210> 1399
<211> 68
<212> PRT
<213> Homo sapiens

<400> 1399

```
Met Gly Ala Val Leu Leu Val Cys Leu Gln Thr Ser Ile Ala Ala Arg
1               5                   10                  15
Asp Asp Leu Lys Asp Ala Val Asp Ser Gly Leu Leu Leu Ala Asn Ser
            20                  25                  30
Leu Ser His Phe Val Pro Leu Val Val Arg Asn Tyr Leu Val His Cys
        35                  40                  45
Asn Leu Leu Gln Thr Leu Lys Phe Leu Leu Gly Asn Cys Thr Ala Gly
    50                  55                  60
Lys Ala Ser  *
65      67
```

<210> 1400
<211> 54
<212> PRT
<213> Homo sapiens

<400> 1400

```
Met Ala Val Ala Phe Val Leu Ser Leu Gly Val Ala Ala Leu Tyr Lys
1               5                   10                  15
Phe Arg Val Ala Asp Gln Arg Lys Lys Ala Tyr Ala Asp Phe Tyr Arg
            20                  25                  30
Asn Tyr Asp Val Met Lys Asp Phe Glu Glu Met Arg Lys Ala Gly Ile
        35                  40                  45
Phe Gln Ser Val Lys  *
    50          53
```

<210> 1401
<211> 232
<212> PRT
<213> Homo sapiens

<400> 1401

```
Met Leu Phe Ala Phe Ile Ser Leu Leu Val Met Leu Pro Thr Trp Trp
1               5                   10                  15
Ile Val Ser Ser Trp Leu Val Trp Gly Val Ile Leu Phe Val Tyr Leu
```

```
            20                       25                       30
Val Ile Arg Ala Leu Arg Leu Trp Arg Thr Ala Lys Leu Gln Val Thr
        35                      40                      45
Leu Lys Lys Tyr Ser Val His Leu Glu Asp Met Ala Thr Asn Ser Arg
    50                      55                      60
Ala Phe Thr Asn Leu Val Arg Lys Ala Leu Arg Leu Ile Gln Glu Thr
65                      70                      75                  80
Glu Val Ile Ser Arg Gly Phe Thr Leu Leu Leu Asp Arg Val Ser Ala
                85                      90                      95
Ala Cys Pro Phe Asn Lys Ala Gly Gln His Pro Ser Gln His Leu Ile
            100                     105                     110
Gly Leu Arg Lys Ala Val Tyr Arg Thr Leu Arg Ala Ser Phe Gln Ala
        115                     120                     125
Ala Arg Leu Ala Thr Leu Tyr Met Leu Lys Asn Tyr Pro Leu Asn Ser
    130                     135                     140
Glu Ser Asp Asn Val Thr Asn Tyr Ile Cys Val Val Pro Phe Lys Glu
145                     150                     155                 160
Leu Gly Leu Gly Leu Ser Glu Glu Gln Ile Ser Glu Glu Glu Ala His
                165                     170                     175
Lys Leu Tyr Arg Trp Leu Gln Pro Ala Cys Ile Glu Gly Phe Val Pro
            180                     185                     190
Thr Leu Gly Gly Thr Glu Phe Arg Val Leu Gln Thr Val Ser Pro Ile
        195                     200                     205
Thr Phe Tyr Ser Gln Phe Thr Ser Trp Ala Leu Thr Tyr Ser Ser Thr
    210                     215                     220
Ser Ala Ser Ser Tyr Leu Ile  *
225                     230 231
```

```
<210> 1402
<211> 48
<212> PRT
<213> Homo sapiens

<400> 1402
Met Ala Pro Ala Arg Pro Trp Trp Leu Thr Pro Val Ile Pro Ala Leu
 1               5                  10                  15
Trp Glu Ala Glu Glu Asp Gly Ser Arg Gly Gln Glu Phe Lys Thr Ser
            20                  25                  30
Leu Ala Ser Met Val Lys Pro Arg Leu Tyr Tyr Lys Tyr Lys Asn  *
        35                  40                  45      47
```

```
<210> 1403
<211> 53
<212> PRT
<213> Homo sapiens

<400> 1403
Met Leu Trp Arg Leu Ile Ile Ile Leu Cys Glu Ala Leu Gln Arg Lys
 1               5                  10                  15
Ser Arg Leu Leu Ala Asp Cys Asp His Phe Ser Phe Pro Asn Arg Tyr
            20                  25                  30
Glu Arg Lys Leu Leu Leu Asp Phe Thr Val Arg Ile Trp Ile Gln Thr
        35                  40                  45
```

Tyr Cys Pro His *
50 52

<210> 1404
<211> 90
<212> PRT
<213> Homo sapiens

<400> 1404
Met Arg Val Phe Cys Val Gly Leu Leu Leu Phe Ser Val Thr Trp Ala
1 5 10 15
Ala Pro Thr Phe Gln Pro Gln Thr Glu Lys Thr Lys Gln Ser Cys Val
20 25 30
Glu Glu Gln Arg Gln Glu Glu Lys Asn Lys Asp Asn Ile Gly Phe His
35 40 45
His Leu Gly Lys Arg Ile Asn Gln Glu Leu Ser Ser Lys Glu Asn Ile
50 55 60
Val Gln Glu Arg Lys Lys Asp Leu Ser Leu Ser Glu Ala Ser Glu Asn
65 70 75 80
Lys Gly Ser Ser Lys Ser Gln Asn Tyr Phe
85 90

<210> 1405
<211> 477
<212> PRT
<213> Homo sapiens

<400> 1405
Met Ala Gly Arg Gly Gly Ser Ala Leu Leu Ala Leu Cys Gly Ala Leu
1 5 10 15
Ala Ala Cys Gly Trp Leu Leu Gly Ala Glu Ala Gln Glu Pro Gly Ala
20 25 30
Pro Ala Ala Gly Met Arg Arg Arg Arg Arg Leu Gln Gln Glu Asp Gly
35 40 45
Ile Ser Phe Glu Tyr His Arg Tyr Pro Glu Leu Arg Glu Ala Leu Val
50 55 60
Ser Val Trp Leu Gln Cys Thr Ala Ile Ser Arg Ile Tyr Thr Val Gly
65 70 75 80
Arg Ser Phe Glu Gly Arg Glu Leu Leu Val Ile Glu Leu Ser Asp Asn
85 90 95
Pro Gly Val His Glu Pro Gly Glu Pro Glu Phe Lys Tyr Ile Gly Asn
100 105 110
Met His Gly Asn Glu Ala Val Gly Arg Glu Leu Leu Ile Phe Leu Ala
115 120 125
Gln Tyr Leu Cys Asn Glu Tyr Gln Lys Gly Asn Glu Thr Ile Val Asn
130 135 140
Leu Ile His Ser Thr Arg Ile His Ile Met Pro Ser Leu Asn Pro Asp
145 150 155 160
Gly Phe Glu Lys Ala Ala Ser Gln Pro Gly Glu Leu Lys Asp Trp Phe
165 170 175
Val Gly Arg Ser Asn Ala Gln Gly Ile Asp Leu Asn Arg Asn Phe Pro
180 185 190
Asp Leu Asp Arg Ile Val Tyr Val Asn Glu Lys Glu Gly Gly Pro Asn

```
        195                 200                 205
Asn His Leu Leu Lys Asn Met Lys Lys Ile Val Asp Gln Asn Thr Lys
    210                 215                 220
Leu Ala Pro Glu Thr Lys Ala Val Ile His Trp Ile Met Asp Ile Pro
225                 230                 235                 240
Phe Val Leu Ser Ala Asn Leu His Gly Gly Asp Leu Val Ala Asn Tyr
                245                 250                 255
Pro Tyr Asp Glu Thr Arg Ser Gly Ser Ala His Glu Tyr Ser Ser Ser
            260                 265                 270
Pro Asp Asp Ala Ile Phe Gln Ser Leu Ala Arg Ala Tyr Ser Ser Phe
        275                 280                 285
Asn Pro Ala Met Ser Asp Pro Asn Arg Pro Pro Cys Arg Lys Asn Asp
    290                 295                 300
Asp Asp Ser Ser Phe Val Asp Gly Thr Thr Asn Gly Gly Ala Trp Tyr
305                 310                 315                 320
Ser Val Pro Gly Gly Met Gln Asp Phe Asn Tyr Leu Ser Ser Asn Cys
                325                 330                 335
Phe Glu Ile Thr Val Glu Leu Ser Cys Glu Lys Phe Pro Pro Glu Glu
            340                 345                 350
Thr Leu Lys Thr Tyr Trp Glu Asp Asn Lys Asn Ser Leu Ile Ser Tyr
        355                 360                 365
Leu Glu Gln Ile His Arg Gly Val Lys Gly Phe Val Arg Asp Leu Gln
    370                 375                 380
Gly Asn Pro Ile Ala Asn Ala Thr Ile Ser Val Glu Gly Ile Asp His
385                 390                 395                 400
Asp Val Thr Ser Ala Lys Asp Gly Asp Tyr Trp Arg Leu Leu Ile Pro
                405                 410                 415
Gly Asn Tyr Lys Leu Thr Ala Ser Ala Pro Gly Tyr Leu Ala Ile Thr
            420                 425                 430
Lys Lys Val Ala Val Pro Tyr Ser Pro Ala Ala Gly Val Asp Phe Glu
        435                 440                 445
Leu Glu Ser Phe Ser Glu Arg Lys Glu Glu Glu Lys Glu Glu Leu Met
    450                 455                 460
Glu Trp Trp Lys Met Met Ser Glu Thr Leu Asn Phe  *
465                 470                 475 476
```

<210> 1406
<211> 55
<212> PRT
<213> Homo sapiens

<400> 1406

```
Met Phe Ile Gly Ile Trp Val Ser Leu Tyr Gln Val Leu Trp Leu Lys
 1               5                  10                  15
Glu Leu Leu Trp Gly His Tyr Ile Phe Trp Val Ser Arg Lys Met Phe
            20                  25                  30
Val Tyr Gly Gly Val Gly Gly Lys Thr Ala Asn Ile Cys Arg Lys Gly
        35                  40                  45
Arg Ile Ile Lys Lys Val  *
    50              54
```

<210> 1407
<211> 66
<212> PRT

<213> Homo sapiens

<400> 1407
Met Leu Leu Gly Val Arg Ala Val Pro Leu Cys Ser Ala Trp Gln Gly
1 5 10 15
Ala Val Gly Leu Val Ser Leu Thr Ile Ser Ile Cys Lys His Gly Leu
20 25 30
Ser Phe Gln Gln Asn Leu Val Pro Gly Lys Ser Asn Val Pro Lys Ala
35 40 45
Ser Asp Met Pro Arg Cys Pro Pro Val Asp Ala Ala Ala Asn Ser Arg
50 55 60
Ser Met
65 66

<210> 1408
<211> 58
<212> PRT
<213> Homo sapiens

<400> 1408
Met Leu Leu Lys Phe Leu Cys Glu Cys Met Pro Ser Leu Leu Leu Ser
1 5 10 15
Glu Phe Leu Asp Ser Pro Arg Ser Gly Ile Asp Gly Ser Asn Gly Asn
20 25 30
Ser Met Phe Asn Phe Val Lys Asn Cys His Phe Pro Thr Ala Ala Ala
35 40 45
Pro Phe Pro Thr Pro Thr Ser Arg Val *
50 55 57

<210> 1409
<211> 72
<212> PRT
<213> Homo sapiens

<400> 1409
Met Ile Glu Thr Trp Leu Trp Leu Leu Leu Leu Asn Val Gly Gly Thr
1 5 10 15
Gly Gln Trp Ser Gly Pro Thr Phe Arg Arg Glu Asn Val Leu Pro Ala
20 25 30
Ala His Ile Gly Pro Lys Tyr Gly Pro Leu Leu Pro Ser Thr Ala Lys
35 40 45
Gly Thr Val Lys Val Ser Cys Pro Ser Ser Thr Pro His Pro Pro Leu
50 55 60
Gln Gly Lys Gly Thr Pro Asp *
65 70 71

<210> 1410
<211> 53
<212> PRT
<213> Homo sapiens

```
     <400> 1410
Met Arg Phe Leu Leu Leu Trp Phe Ile Leu Arg Gly Arg Gln Leu Val
 1               5                  10                  15
Pro Leu Arg Pro Arg Arg Ser Pro Leu Pro Asp Thr Asn Ala Pro Leu
            20                  25                  30
Pro Gly Leu Gly Gly Gly Asp Gly Ser Thr Gln Thr Pro Phe Ala Gln
        35                  40                  45
Ser Arg Arg Leu  *
    50      52


     <210> 1411
     <211> 82
     <212> PRT
     <213> Homo sapiens


     <400> 1411
Met Ala Ser Gln Ser Met Cys Phe Leu Trp Leu Ala Pro Val Thr Trp
 1               5                  10                  15
Cys Val Met Phe Ser Ser Arg Thr Cys Tyr Ser Pro Cys Gly Asn Phe
            20                  25                  30
Ser Thr Ala Pro Gly Arg Val Ile Phe His Ser Trp Asp Arg Ala Gln
        35                  40                  45
Phe Val Tyr Ser Phe Leu Ser Arg Trp Arg Leu Gly Leu Phe Pro Pro
    50                  55                  60
Leu Ala Ser Val Asn Gly Asp Ala Val Ile Met Gly Val Pro Val Phe
65                  70                  75                  80
Val  *
81


     <210> 1412
     <211> 72
     <212> PRT
     <213> Homo sapiens


     <400> 1412
Met Phe Leu Leu Leu Phe Cys Leu Met Phe Asp Phe Thr Lys Val Phe
 1               5                  10                  15
Phe Ile Leu Leu Leu His Ile Phe Cys Leu Ser Thr Cys Leu Phe Leu
            20                  25                  30
Gly Leu His Ile Cys Ala Ser Phe His Ala Arg Ala Leu Leu Glu Thr
        35                  40                  45
Ala Leu Ile Leu Leu Arg Met Lys Ile Ala Gly Phe Gln Val Ile Leu
    50                  55                  60
Phe Pro Gln Asp Phe Val Leu  *
65                  70  71


     <210> 1413
     <211> 59
     <212> PRT
```

<213> Homo sapiens

<400> 1413
```
Met Met Thr Ile Lys Glu Phe Thr Leu Leu Leu Val Ser Leu Gln Phe
 1               5                  10                  15
Ser Thr Phe Pro Ser Lys Lys Phe Leu Leu Glu Thr His Phe Leu Lys
            20                  25                  30
Asn Ser Glu Asn Trp Leu Gly Val Val Ala His Ala Cys Ser Leu Ser
        35                  40                  45
Thr Leu Gly Trp Pro Arg Arg Arg Thr Ala  *
    50                  55          58
```

<210> 1414
<211> 78
<212> PRT
<213> Homo sapiens

<400> 1414
```
Met Leu Arg Leu Asp Ile Ile Asn Ser Leu Val Thr Thr Val Phe Met
 1               5                  10                  15
Leu Ile Val Ser Val Leu Ala Leu Ile Pro Glu Thr Thr Thr Leu Thr
            20                  25                  30
Val Gly Gly Gly Val Phe Ala Leu Val Thr Ala Val Cys Cys Leu Ala
        35                  40                  45
Asp Gly Ala Leu Ile Tyr Arg Lys Leu Leu Phe Asn Pro Ser Gly Pro
    50                  55                  60
Tyr Gln Lys Lys Pro Val His Glu Lys Lys Glu Val Leu  *
65                  70                  75      77
```

<210> 1415
<211> 171
<212> PRT
<213> Homo sapiens

<400> 1415
```
Met His Met Met Lys Leu Ser Ile Lys Val Leu Leu Gln Ser Ala Leu
 1               5                  10                  15
Ser Leu Gly Arg Ser Leu Asp Ala Asp His Ala Pro Leu Gln Gln Phe
            20                  25                  30
Phe Val Val Met Glu His Cys Leu Lys His Gly Leu Lys Val Lys Lys
        35                  40                  45
Ser Phe Ile Gly Gln Asn Lys Ser Phe Phe Gly Pro Leu Glu Leu Val
    50                  55                  60
Glu Lys Leu Cys Pro Glu Ala Ser Asp Ile Ala Thr Ser Val Arg Asn
65                  70                  75                  80
Leu Pro Glu Leu Lys Thr Ala Val Gly Arg Gly Arg Ala Trp Leu Tyr
                85                  90                  95
Leu Ala Leu Met Gln Lys Lys Leu Ala Asp Tyr Leu Lys Val Leu Ile
            100                 105                 110
Asp Asn Lys His Leu Leu Ser Glu Phe Tyr Glu Pro Glu Ala Leu Met
        115                 120                 125
Met Glu Glu Glu Gly Met Val Ile Val Gly Leu Leu Val Gly Leu Asn
```

```
    130                 135                 140
Val Leu Asp Ala Asn Leu Trp Leu Glu Arg Arg Arg Leu Gly Phe Ser
145                 150                 155                 160
Gly Trp Ser Asn Arg Phe Phe Pro Leu Pro  *
                165                 170
```

<210> 1416
<211> 77
<212> PRT
<213> Homo sapiens

<400> 1416

```
Met Leu Thr Arg Leu Val Leu Ser Ala His Leu Ser Ser Thr Thr Phe
 1               5                  10                  15
Pro Pro Trp Thr His Ala Ala Ile Ser Trp Glu Leu Asp Asn Val Leu
            20                  25                  30
Met Pro Ser Pro Arg Ile Trp Pro Gln Val Thr Pro Thr Ala Gly Gln
        35                  40                  45
Asp Val His Ala Ile Val Thr Arg Thr Cys Glu Ser Val Leu Ser Ser
    50                  55                  60
Val Val Tyr Thr His Gly Cys Gly Cys Val Arg Cys  *
 65                  70                  75  76
```

<210> 1417
<211> 249
<212> PRT
<213> Homo sapiens

<400> 1417

```
Met Glu Lys Ile Pro Glu Ile Gly Lys Phe Gly Glu Lys Ala Pro Pro
 1               5                  10                  15
Ala Pro Ser His Val Trp Arg Pro Ala Ala Leu Phe Leu Thr Leu Leu
            20                  25                  30
Cys Leu Leu Leu Leu Ile Gly Leu Gly Val Leu Ala Ser Met Phe His
        35                  40                  45
Val Thr Leu Lys Ile Glu Met Lys Lys Met Asn Lys Leu Gln Asn Ile
    50                  55                  60
Ser Glu Glu Leu Gln Arg Asn Ile Ser Leu Gln Leu Met Ser Asn Met
 65                  70                  75                  80
Asn Ile Ser Asn Lys Ile Arg Asn Leu Ser Thr Thr Leu Gln Thr Ile
                85                  90                  95
Ala Thr Lys Leu Cys Arg Glu Leu Tyr Ser Lys Glu Gln Glu His Lys
           100                 105                 110
Cys Lys Pro Cys Pro Arg Arg Trp Ile Trp His Lys Asp Ser Cys Tyr
       115                 120                 125
Phe Leu Ser Asp Asp Val Gln Thr Trp Gln Glu Ser Lys Met Ala Cys
    130                 135                 140
Ala Ala Gln Asn Ala Ser Leu Leu Lys Ile Asn Asn Lys Asn Ala Leu
145                 150                 155                 160
Glu Phe Ile Lys Ser Gln Ser Arg Ser Tyr Asp Tyr Trp Leu Gly Leu
                165                 170                 175
Ser Pro Glu Glu Asp Ser Thr Arg Gly Met Arg Val Asp Asn Ile Ile
           180                 185                 190
```

Asn Ser Ser Ala Trp Val Ile Arg Asn Ala Pro Asp Leu Asn Asn Met
195 200 205
Tyr Cys Gly Tyr Ile Asn Arg Leu Tyr Val Gln Tyr Tyr His Cys Thr
210 215 220
Tyr Lys Gln Arg Met Ile Cys Glu Lys Met Ala Asn Pro Val Gln Leu
225 230 235 240
Gly Ser Thr Tyr Phe Arg Glu Ala *
245 248

<210> 1418
<211> 65
<212> PRT
<213> Homo sapiens

<400> 1418
Met Gly Leu Lys Asn Val Phe Leu Pro Val Phe Leu Pro Phe Leu Leu
1 5 10 15
Tyr Ser Glu Phe Leu Ser Leu Pro Pro Ser Leu Ser Ser Ser Leu Leu
20 25 30
Pro Phe Leu Pro Phe Ser Leu Pro Gly His Phe Ser Asn Leu His Gln
35 40 45
Arg Tyr Leu Lys Cys Trp Tyr Leu Arg Ile Ser Val Thr Pro Leu Ile
50 55 60 64
*

<210> 1419
<211> 468
<212> PRT
<213> Homo sapiens

<400> 1419
Met Leu Leu Leu Leu Leu Leu Pro Leu Leu Trp Gly Arg Glu Arg Val
1 5 10 15
Glu Gly Gln Lys Ser Asn Arg Lys Asp Tyr Ser Leu Thr Met Gln Ser
20 25 30
Ser Val Thr Val Gln Glu Gly Met Cys Val His Val Arg Cys Ser Phe
35 40 45
Ser Tyr Pro Val Asp Ser Gln Thr Asp Ser Asp Pro Val His Gly Tyr
50 55 60
Trp Phe Arg Ala Gly Asn Asp Ile Ser Trp Lys Ala Pro Val Ala Thr
65 70 75 80
Asn Asn Pro Ala Trp Ala Val Gln Glu Glu Thr Arg Asp Arg Phe His
85 90 95
Leu Leu Gly Asp Pro Gln Thr Lys Asn Cys Thr Leu Ser Ile Arg Asp
100 105 110
Ala Arg Met Ser Asp Ala Gly Arg Tyr Phe Phe Arg Met Glu Lys Gly
115 120 125
Asn Ile Lys Trp Asn Tyr Lys Tyr Asp Gln Leu Ser Val Asn Val Thr
130 135 140
Ala Leu Thr His Arg Pro Asn Ile Leu Ile Pro Gly Thr Leu Glu Ser
145 150 155 160
Gly Cys Phe Gln Asn Leu Thr Cys Ser Val Pro Trp Ala Cys Glu Gln

```
                165                 170                 175
Gly Thr Pro Pro Met Ile Ser Trp Met Gly Thr Ser Val Ser Pro Leu
            180                 185                 190
His Pro Ser Thr Thr Arg Ser Ser Val Leu Thr Leu Ile Pro Gln Pro
        195                 200                 205
Gln His His Gly Thr Ser Leu Thr Cys Gln Val Thr Leu Pro Gly Ala
    210                 215                 220
Gly Val Thr Thr Asn Arg Thr Ile Gln Leu Asn Val Ser Tyr Pro Pro
225                 230                 235                 240
Gln Asn Leu Thr Val Thr Val Phe Gln Gly Glu Gly Thr Ala Ser Thr
                245                 250                 255
Ala Leu Gly Asn Ser Ser Ser Leu Ser Val Leu Glu Gly Gln Ser Leu
            260                 265                 270
Arg Leu Val Cys Ala Val Asp Ser Asn Pro Pro Ala Arg Leu Ser Trp
        275                 280                 285
Thr Trp Arg Ser Leu Thr Leu Tyr Pro Ser Gln Pro Ser Asn Pro Leu
    290                 295                 300
Val Leu Glu Leu Gln Val His Leu Gly Asp Glu Gly Glu Phe Thr Cys
305                 310                 315                 320
Arg Ala Gln Asn Ser Leu Gly Ser Gln His Val Ser Leu Asn Leu Ser
                325                 330                 335
Leu Gln Gln Glu Tyr Thr Gly Lys Met Arg Pro Val Ser Gly Val Leu
            340                 345                 350
Leu Gly Ala Val Gly Gly Ala Gly Ala Thr Ala Leu Val Phe Leu Ser
        355                 360                 365
Phe Cys Val Ile Phe Ile Val Val Arg Ser Cys Arg Lys Lys Ser Ala
    370                 375                 380
Arg Pro Ala Ala Asp Val Gly Asp Ile Gly Met Lys Asp Ala Asn Thr
385                 390                 395                 400
Ile Arg Gly Ser Ala Ser Gln Gly Asn Leu Thr Glu Ser Trp Ala Asp
                405                 410                 415
Asp Asn Pro Arg His His Gly Leu Ala Ala His Ser Ser Gly Glu Glu
            420                 425                 430
Arg Glu Ile Gln Tyr Ala Pro Leu Ser Phe His Lys Gly Glu Pro Gln
        435                 440                 445
Asp Leu Ser Gly Gln Glu Ala Thr Asn Asn Glu Tyr Ser Glu Ile Lys
    450                 455                 460
Ile Pro Lys  *
465     467


     <210> 1420
     <211> 150
     <212> PRT
     <213> Homo sapiens


     <400> 1420
Met Ile Arg Cys Leu Ala Gln Pro Ala Ala Val Leu Ser Ser Leu Gly
 1               5                  10                  15
Leu Ala Gln Val Leu Gly Asp Ser Gly Arg Asp Glu Gln Val Leu Leu
            20                  25                  30
Arg Arg Ser Phe Arg Ala Glu Gly Cys Val Leu Cys Leu Cys Thr Trp
        35                  40                  45
Gly Thr Ala Val Pro Trp His Lys Val Glu Gly Ser Gly Gly Pro Cys
    50                  55                  60
Arg Ser Ala Ala Pro Leu Pro Ala Ser Ala Pro Phe Ser Ile Asp Gly
65                  70                  75                  80
```

Arg Ala Val Pro Trp Val Phe Ser Ala Leu Gln Ala Glu Val Gly Val
85 90 95
Leu Gly Glu Gln Met Arg Asp Gly Arg Gly Leu Cys Gly Ser His Pro
100 105 110
Trp Val Leu Gln Leu Ser Trp Pro Gly Val Phe Pro Gln Cys Trp Leu
115 120 125
Cys Pro Arg Leu Val Cys Leu Ala Lys Gln Asn Trp Gln Cys Pro Phe
130 135 140
Glu Thr Pro Arg Lys *
145 149

<210> 1421
<211> 89
<212> PRT
<213> Homo sapiens

<400> 1421
Met Tyr Val Phe Leu Leu Cys Pro Ala Cys Gly Arg Leu Met Gly Ser
1 5 10 15
Thr Tyr Met Arg Leu Leu Pro Gln Ser Glu Pro Ala Leu His Asn Arg
20 25 30
Ile Leu Arg Gln Thr Glu Pro Leu Leu Tyr Phe Lys Arg Gly Lys Gln
35 40 45
Gln Gly Leu Phe Tyr Ala Ser Phe Pro Ala Val His Arg Met Asp Ser
50 55 60
Leu Leu Arg Arg Thr Val Val Ile Leu Tyr Lys Arg Thr Asn Thr Val
65 70 75 80
Gly Val Ser Leu Phe Gln Asn Ala *
85 88

<210> 1422
<211> 83
<212> PRT
<213> Homo sapiens

<400> 1422
Met Met Thr Trp Ala Ser Leu Ala Leu Gly Leu Thr Arg Ala Leu Gly
1 5 10 15
Gly Met Gly Ser Phe Leu Leu Arg Ile Leu Gly Trp Ser Trp Ala Met
20 25 30
Gly Ser Arg Ser Arg Ala Arg Trp Pro Arg Gly Arg Leu Gly Phe Thr
35 40 45
Ser Met Leu Ser Cys Met Arg Gln Cys Ser Val Cys Arg Met Ile Met
50 55 60
Ser Leu Val Glu Val Leu Val Ala Thr Ser Gln Val Val Lys Leu Trp
65 70 75 80
Ser Arg *
82

<210> 1423
<211> 54

<212> PRT
<213> Homo sapiens

<400> 1423
Met Ile Leu Phe Pro Leu Cys Pro Ser Ile Leu Ser Leu Lys Pro Lys
1 5 10 15
Lys Lys Glu Ala Leu Pro Ser Leu Ser Val Met Gly Thr Val Phe Leu
20 25 30
Leu Val Ser Cys Ser Leu Pro Ser Pro Ala Ala Cys Gly Arg Asn Ala
35 40 45
Ala Thr Ala Gln His *
50 53

<210> 1424
<211> 73
<212> PRT
<213> Homo sapiens

<400> 1424
Met Cys Phe Ser Cys Leu Pro Leu Gln Cys Leu Ala Met Gly His Lys
1 5 10 15
His Tyr Pro Ala Val Gly Arg Leu Ala Lys Arg Ser Gln Leu Ala Ser
20 25 30
Pro Ala Ser Ser Arg Glu Trp Asn His Gly Ser Asn Thr Leu Leu Arg
35 40 45
Lys Gln Lys Leu Tyr Gly His Ile Phe His Leu Leu Ser Pro Arg Asn
50 55 60
His Met Tyr Cys Asp Pro Ala His *
65 70 72

<210> 1425
<211> 245
<212> PRT
<213> Homo sapiens

<400> 1425
Met Ala Cys Tyr Leu Leu Val Ala Asn Ile Leu Leu Val Asn Leu Leu
1 5 10 15
Ile Ala Val Phe Asn Asn Thr Phe Phe Glu Val Lys Ser Ile Ser Asn
20 25 30
Gln Val Trp Lys Phe Gln Arg Tyr Gln Leu Ile Met Thr Phe His Glu
35 40 45
Arg Pro Val Leu Pro Pro Pro Leu Ile Ile Phe Ser His Met Thr Met
50 55 60
Ile Phe Gln His Leu Cys Cys Arg Trp Arg Lys His Glu Ser Asp Pro
65 70 75 80
Asp Glu Arg Asp Tyr Gly Leu Lys Leu Phe Ile Thr Asp Asp Glu Leu
85 90 95
Lys Lys Val His Asp Phe Glu Glu Gln Cys Ile Glu Glu Tyr Phe Arg
100 105 110
Glu Lys Asp Asp Arg Phe Asn Ser Ser Asn Asp Glu Arg Ile Arg Val
115 120 125

Thr Ser Glu Arg Val Glu Asn Met Ser Met Arg Leu Glu Glu Val Asn
    130                 135                 140
Glu Arg Glu His Ser Met Lys Ala Ser Leu Gln Thr Val Asp Ile Arg
145                 150                 155                 160
Leu Ala Gln Leu Glu Asp Leu Ile Gly Arg Met Ala Thr Ala Leu Glu
                165                 170                 175
Arg Leu Thr Gly Leu Glu Arg Ala Glu Ser Asn Lys Ile Arg Ser Arg
            180                 185                 190
Thr Ser Ser Asp Cys Thr Asp Ala Arg Leu His Trp Pro Val Arg Ala
        195                 200                 205
Ala Leu Thr Ser Gln Glu Arg Glu His Leu Ser Ala Pro Lys Arg Gly
    210                 215                 220
Leu Glu Pro Trp Gln Asn Ile Leu Phe Ile Gln Tyr Lys Pro Ala Ala
225                 230                 235                 240
Ser Ser Ser Thr  *
            244

<210> 1426
<211> 520
<212> PRT
<213> Homo sapiens

<221> misc_feature
<222> (1)...(520)
<223> Xaa = any amino acid or nothing

<400> 1426
Met Asp Ile Leu Leu Leu Leu Leu Phe Phe Met Ile Ile Phe Ala Ile
  1               5                  10                  15
Leu Gly Phe Tyr Leu Phe Ser Pro Asn Pro Ser Asp Pro Tyr Phe Ser
            20                  25                  30
Thr Leu Glu Asn Ser Ile Val Ser Leu Phe Val Leu Leu Thr Thr Ala
        35                  40                  45
Asn Phe Pro Asp Val Met Met Pro Ser Tyr Ser Arg Asn Pro Trp Ser
    50                  55                  60
Cys Val Phe Phe Ile Val Tyr Leu Ser Ile Glu Leu Tyr Phe Ile Met
65                  70                  75                  80
Asn Leu Leu Leu Ala Val Val Phe Asp Thr Phe Asn Asp Ile Glu Lys
                85                  90                  95
Arg Lys Phe Lys Ser Leu Leu Leu His Lys Arg Thr Ala Ile Gln His
            100                 105                 110
Ala Tyr Arg Leu Leu Ile Ser Gln Arg Arg Pro Ala Gly Ile Ser Tyr
        115                 120                 125
Arg Gln Phe Glu Gly Leu Met Arg Phe Tyr Lys Pro Arg Met Ser Ala
    130                 135                 140
Arg Glu Arg Tyr Leu Thr Phe Lys Ala Leu Asn Gln Asn Asn Thr Pro
145                 150                 155                 160
Leu Leu Ser Leu Lys Asp Phe Tyr Asp Ile Tyr Glu Val Ala Ala Leu
                165                 170                 175
Lys Trp Lys Ala Thr Lys Asn Arg Glu His Trp Val Asp Glu Leu Pro
            180                 185                 190
Arg Thr Ala Leu Leu Ile Phe Lys Gly Ile Asn Ile Leu Val Lys Ala
        195                 200                 205
Lys Ala Phe Gln Tyr Phe Met Tyr Leu Val Val Ala Val Asn Gly Val
    210                 215                 220
Trp Ile Leu Val Glu Thr Phe Met Leu Lys Gly Gly Asn Phe Phe Ser

```
225                 230                 235                 240
Lys His Val Pro Trp Ser Tyr Leu Val Phe Leu Thr Ile Tyr Gly Val
                245                 250                 255
Glu Leu Phe Leu Lys Val Ala Gly Leu Gly Pro Val Glu Tyr Leu Ser
            260                 265                 270
Ser Gly Trp Asn Leu Phe Asp Phe Ser Val Thr Val Phe Ala Phe Leu
        275                 280                 285
Gly Leu Leu Ala Leu Ala Leu Asn Met Glu Pro Phe Tyr Phe Ile Val
    290                 295                 300
Val Leu Arg Pro Leu Gln Leu Leu Arg Leu Phe Lys Leu Lys Glu Arg
305                 310                 315                 320
Tyr Arg Asn Val Leu Asp Thr Met Phe Glu Leu Leu Pro Arg Met Ala
                325                 330                 335
Ser Leu Gly Leu Thr Leu Leu Ile Phe Tyr Tyr Ser Phe Ala Ile Val
            340                 345                 350
Gly Met Glu Phe Phe Cys Gly Ile Val Phe Pro Asn Cys Cys Asn Thr
        355                 360                 365
Ser Thr Val Ala Asp Ala Tyr Arg Trp Arg Asn His Thr Val Gly Asn
    370                 375                 380
Arg Thr Val Val Glu Glu Gly Tyr Tyr Tyr Leu Asn Asn Phe Asp Asn
385                 390                 395                 400
Ile Leu Asn Ser Phe Val Thr Leu Phe Glu Leu Thr Val Val Asn Asn
                405                 410                 415
Trp Tyr Ile Ile Met Glu Gly Val Thr Ser Gln Thr Ser His Trp Ser
            420                 425                 430
Arg Leu Tyr Phe Met Thr Phe Tyr Ile Ala Thr Met Val Val Met Thr
        435                 440                 445
Ile Ile Val Ala Phe Ile Leu Glu Ala Phe Val Phe Arg Met Asn Tyr
    450                 455                 460
Ser Arg Lys Asn Gln Asp Ser Glu Val Asp Gly Gly Ile Thr Leu Glu
465                 470                 475                 480
Lys Glu Ile Ser Lys Glu Glu Leu Val Ala Val Leu Glu Leu Tyr Arg
                485                 490                 495
Glu Ala Arg Xaa Ala Ser Ser Asp Val Thr Arg Leu Leu Glu Thr Leu
            500                 505                 510
Ser Gln Met Glu Arg Tyr Gln Gln
        515                 520
```

<210> 1427
<211> 106
<212> PRT
<213> Homo sapiens

<400> 1427

```
Met Ser Pro Gln His Leu Leu Leu Thr Leu Pro Leu Pro Leu Arg Ser
 1               5                  10                  15
Pro Ile Leu Phe Ser His Thr Ala Gln Leu Leu Val Leu Thr Arg Ile
            20                  25                  30
Ala Phe Arg Ala Cys Glu Leu Phe Phe Phe Val Met Val Ser Leu Cys
        35                  40                  45
Cys Pro Gly Ile His Ser Phe Ile Ala Thr Ile Thr Tyr Glu Arg Asn
    50                  55                  60
Ala Phe Gln Ser Ile Ser Ser Val Gln Gln Gln His Leu His Phe Gly
65                  70                  75                  80
Cys Ala Leu Ser Pro Pro Ala Pro Arg Glu Ser Phe Ser Pro Cys Leu
                85                  90                  95
```

Thr Thr His Arg Leu Pro Ser Cys Phe *
100 105

<210> 1428
<211> 841
<212> PRT
<213> Homo sapiens

<400> 1428
Met Ala Leu Ala Ser Ala Ala Pro Gly Ser Ile Phe Cys Lys Gln Leu
1 5 10 15
Leu Phe Ser Leu Leu Val Leu Thr Leu Leu Cys Asp Ala Cys Gln Lys
20 25 30
Val Tyr Leu Arg Val Pro Ser His Leu Gln Ala Glu Thr Leu Val Gly
35 40 45
Lys Val Asn Leu Glu Glu Cys Leu Lys Ser Ala Ser Leu Ile Arg Ser
50 55 60
Ser Asp Pro Ala Phe Arg Ile Leu Glu Asp Gly Ser Ile Tyr Thr Thr
65 70 75 80
His Asp Leu Ile Leu Ser Ser Glu Arg Lys Ser Phe Ser Ile Phe Leu
85 90 95
Ser Asp Gly Gln Arg Arg Glu Gln Gln Glu Ile Lys Val Val Leu Ser
100 105 110
Ala Arg Glu Asn Lys Ser Pro Lys Lys Arg His Thr Lys Asp Thr Ala
115 120 125
Leu Lys Arg Ser Lys Arg Arg Trp Ala Pro Ile Pro Ala Ser Leu Met
130 135 140
Glu Asn Ser Leu Gly Pro Phe Pro Gln His Val Gln Gln Ile Gln Ser
145 150 155 160
Asp Ala Ala Gln Asn Tyr Thr Ile Phe Tyr Ser Ile Ser Gly Pro Gly
165 170 175
Val Asp Lys Glu Pro Phe Asn Leu Phe Tyr Ile Glu Lys Asp Thr Gly
180 185 190
Asp Ile Phe Cys Thr Arg Ser Ile Asp Arg Glu Lys Tyr Glu Gln Phe
195 200 205
Ala Leu Tyr Gly Tyr Ala Thr Thr Ala Asp Gly Tyr Ala Pro Glu Tyr
210 215 220
Pro Leu Pro Leu Ile Ile Lys Ile Glu Asp Asp Asn Asp Asn Ala Pro
225 230 235 240
Tyr Phe Glu His Arg Val Thr Ile Phe Thr Val Pro Glu Asn Cys Arg
245 250 255
Ser Gly Thr Ser Val Gly Lys Val Thr Ala Thr Asp Leu Asp Glu Pro
260 265 270
Asp Thr Leu His Thr Arg Leu Lys Tyr Lys Ile Leu Gln Gln Ile Pro
275 280 285
Asp His Pro Lys His Phe Ser Ile His Pro Asp Thr Gly Val Ile Thr
290 295 300
Thr Thr Thr Pro Phe Leu Asp Arg Glu Lys Cys Asp Thr Tyr Gln Leu
305 310 315 320
Ile Met Glu Val Arg Asp Met Gly Gly Gln Pro Phe Gly Leu Phe Asn
325 330 335
Thr Gly Thr Ile Thr Ile Ser Leu Glu Asp Glu Asn Asp Asn Pro Pro
340 345 350
Ser Phe Thr Glu Thr Ser Tyr Val Thr Glu Val Glu Glu Asn Arg Ile
355 360 365
Asp Val Glu Ile Leu Arg Met Lys Val Gln Asp Gln Asp Leu Pro Asn

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    370                 375                 380
Thr Pro His Ser Lys Ala Val Tyr Lys Ile Leu Gln Gly Asn Glu Asn
385                 390                 395                 400
Gly Asn Phe Ile Ile Ser Thr Asp Pro Asn Thr Asn Glu Gly Val Leu
                405                 410                 415
Cys Val Val Lys Pro Leu Asn Tyr Glu Val Asn Arg Gln Val Ile Leu
            420                 425                 430
Gln Val Gly Val Ile Asn Glu Ala Gln Phe Ser Lys Ala Ala Ser Ser
        435                 440                 445
Gln Thr Pro Thr Met Cys Thr Thr Thr Val Thr Val Lys Ile Ile Asp
    450                 455                 460
Ser Asp Glu Gly Pro Glu Cys His Pro Pro Val Lys Val Ile Gln Ser
465                 470                 475                 480
Gln Asp Gly Phe Pro Ala Gly Gln Glu Leu Leu Gly Tyr Lys Ala Leu
                485                 490                 495
Asp Pro Glu Ile Ser Ser Gly Glu Gly Leu Arg Tyr Gln Lys Leu Gly
            500                 505                 510
Asp Glu Asp Asn Trp Phe Glu Ile Asn Gln His Thr Gly Asp Leu Arg
        515                 520                 525
Thr Leu Lys Val Leu Asp Arg Glu Ser Lys Phe Val Lys Asn Asn Gln
    530                 535                 540
Tyr Asn Ile Ser Val Val Ala Gly Asp Ala Val Gly Arg Ser Cys Thr
545                 550                 555                 560
Gly Thr Leu Val Val His Leu Asp Asp Tyr Asn Asp His Ala Pro Gln
                565                 570                 575
Ile Asp Lys Glu Val Thr Ile Cys Gln Asn Asn Glu Asp Phe Val Val
            580                 585                 590
Leu Lys Pro Val Asp Pro Asp Gly Pro Glu Asn Gly Pro Pro Phe Gln
        595                 600                 605
Phe Phe Leu Asp Asn Ser Ala Ser Lys Asn Trp Asn Ile Lys Lys Lys
    610                 615                 620
Asp Gly Lys Thr Ala Ile Leu Arg Gln Arg Gln Asn Leu Asp Tyr Asn
625                 630                 635                 640
Tyr Tyr Ser Val Pro Ile Gln Ile Lys Asp Arg His Gly Leu Val Ala
                645                 650                 655
Thr His Met Leu Thr Val Arg Val Cys Asp Cys Ser Thr Pro Ser Glu
            660                 665                 670
Cys Thr Met Lys Asp Lys Ser Thr Arg Asp Val Arg Pro Asn Val Ile
        675                 680                 685
Leu Gly Arg Trp Ala Ile Leu Ala Met Val Leu Gly Ser Val Leu Leu
    690                 695                 700
Leu Cys Ile Leu Phe Thr Cys Phe Cys Val Thr Ala Lys Arg Thr Val
705                 710                 715                 720
Lys Lys Cys Phe Pro Glu Asp Ile Ala Gln Gln Asn Leu Ile Val Ser
                725                 730                 735
Asn Thr Glu Gly Pro Gly Glu Glu Val Thr Glu Ala Asn Ile Arg Leu
            740                 745                 750
Pro Met Gln Thr Ser Asn Ile Cys Asp Thr Ser Met Ser Val Gly Thr
        755                 760                 765
Val Gly Gly Gln Gly Ile Lys Thr Gln Gln Ser Phe Glu Met Val Lys
    770                 775                 780
Gly Gly Tyr Thr Leu Asp Ser Asn Lys Gly Gly Gly His Gln Thr Leu
785                 790                 795                 800
Glu Ser Val Lys Gly Val Gly Gln Gly Asp Thr Gly Arg Tyr Ala Tyr
                805                 810                 815
Thr Asp Trp Gln Ser Phe Thr Gln Pro Arg Leu Gly Glu Glu Ser Ile
            820                 825                 830
Arg Gly His Thr Leu Ile Lys Asn  *
        835                 840
```

<210> 1429
<211> 262
<212> PRT
<213> Homo sapiens

<400> 1429
Met Glu Leu Leu Gln Val Thr Ile Leu Phe Leu Leu Pro Ser Ile Cys
1 5 10 15
Ser Ser Asn Ser Thr Gly Val Leu Glu Ala Ala Asn Asn Ser Leu Val
20 25 30
Val Thr Thr Thr Lys Pro Ser Ile Thr Thr Pro Asn Thr Glu Ser Leu
35 40 45
Gln Lys Asn Val Val Thr Pro Thr Thr Gly Thr Thr Pro Lys Gly Thr
50 55 60
Ile Thr Asn Glu Leu Leu Lys Met Ser Leu Met Ser Thr Ala Thr Phe
65 70 75 80
Leu Thr Ser Lys Asp Glu Gly Leu Lys Ala Thr Thr Thr Asp Val Arg
85 90 95
Lys Asn Asp Ser Ile Ile Ser Asn Val Thr Val Thr Ser Val Thr Leu
100 105 110
Pro Asn Ala Val Ser Thr Leu Gln Ser Ser Lys Pro Lys Thr Glu Thr
115 120 125
Gln Ser Ser Ile Lys Thr Thr Glu Ile Pro Gly Ser Val Leu Gln Pro
130 135 140
Asp Ala Ser Pro Ser Lys Thr Gly Thr Leu Thr Ser Ile Pro Val Thr
145 150 155 160
Ile Pro Glu Asn Thr Ser Gln Ser Gln Val Ile Gly Thr Glu Gly Gly
165 170 175
Lys Asn Ala Ser Thr Ser Ala Thr Ser Arg Ser Tyr Ser Ser Ile Ile
180 185 190
Leu Pro Val Val Ile Ala Leu Ile Val Ile Thr Leu Ser Val Phe Val
195 200 205
Leu Val Gly Leu Tyr Arg Met Cys Trp Lys Ala Asp Pro Gly Thr Pro
210 215 220
Glu Asn Gly Asn Asp Gln Pro Gln Ser Asp Lys Glu Ser Val Lys Leu
225 230 235 240
Leu Thr Val Lys Thr Ile Ser His Glu Ser Gly Glu His Ser Ala Gln
245 250 255
Gly Lys Thr Lys Asn *
260 261

<210> 1430
<211> 66
<212> PRT
<213> Homo sapiens

<400> 1430
Met Ser Tyr Thr Ala Phe Leu Ser Val Cys Cys Leu Pro Leu Leu Pro
1 5 10 15
Leu Cys Asp Phe Ala Leu Tyr Val Leu Leu Asp Lys Phe Lys Gly Gly
20 25 30
Phe Arg Gln Gln Asn Ser Pro Gln Ser Ile Tyr Gln His Asn Pro Tyr

```
        35                  40                  45
Gln Asn Pro Asn Asn Val Leu Ile Phe Leu Gln Lys Trp Lys Asn Arg
    50                  55                  60
Cys  *
 65
```

<210> 1431
<211> 437
<212> PRT
<213> Homo sapiens

<400> 1431

```
Met Leu Lys Val Ser Ala Val Leu Cys Val Cys Ala Ala Ala Trp Cys
  1               5                  10                  15
Ser Gln Ser Leu Ala Ala Ala Ala Ala Val Ala Ala Ala Gly Gly Arg
             20                  25                  30
Ser Asp Gly Gly Asn Phe Leu Asp Asp Lys Gln Trp Leu Thr Thr Ile
         35                  40                  45
Ser Gln Tyr Asp Lys Glu Val Gly Gln Trp Asn Lys Phe Arg Asp Glu
     50                  55                  60
Val Glu Asp Asp Tyr Phe Arg Thr Trp Ser Pro Gly Lys Pro Phe Asp
 65                  70                  75                  80
Gln Ala Leu Asp Pro Ala Lys Asp Pro Cys Leu Lys Met Lys Cys Ser
                 85                  90                  95
Arg His Lys Val Cys Ile Ala Gln Asp Ser Gln Thr Ala Val Cys Ile
            100                 105                 110
Ser His Arg Arg Leu Thr His Arg Met Lys Glu Ala Gly Val Asp His
        115                 120                 125
Arg Gln Trp Arg Gly Pro Ile Leu Ser Thr Cys Lys Gln Cys Pro Val
    130                 135                 140
Val Tyr Pro Ser Pro Val Cys Gly Ser Asp Gly His Thr Tyr Ser Phe
145                 150                 155                 160
Gln Cys Lys Leu Glu Tyr Gln Ala Cys Val Leu Gly Lys Gln Ile Ser
                165                 170                 175
Val Lys Cys Glu Gly His Cys Pro Cys Pro Ser Asp Lys Pro Thr Ser
            180                 185                 190
Thr Ser Arg Asn Val Lys Arg Ala Cys Ser Asp Leu Glu Phe Arg Glu
        195                 200                 205
Val Ala Asn Arg Leu Arg Asp Trp Phe Lys Ala Leu His Glu Ser Gly
    210                 215                 220
Ser Gln Asn Lys Lys Thr Lys Thr Leu Leu Arg Pro Glu Arg Ser Arg
225                 230                 235                 240
Phe Asp Thr Ser Ile Leu Pro Ile Cys Lys Asp Ser Leu Gly Trp Met
                245                 250                 255
Phe Asn Arg Leu Asp Thr Asn Tyr Asp Leu Leu Leu Asp Gln Ser Glu
            260                 265                 270
Leu Arg Ser Ile Tyr Leu Asp Lys Asn Glu Gln Cys Thr Lys Ala Phe
        275                 280                 285
Phe Asn Ser Cys Asp Thr Tyr Lys Asp Ser Leu Ile Ser Asn Asn Glu
    290                 295                 300
Trp Cys Tyr Cys Phe Gln Arg Gln Gln Asp Pro Pro Cys Gln Thr Glu
305                 310                 315                 320
Leu Ser Asn Ile Gln Lys Arg Gln Gly Val Lys Lys Leu Leu Gly Gln
                325                 330                 335
Tyr Ile Pro Leu Cys Asp Glu Asp Gly Tyr Tyr Lys Pro Thr Gln Cys
            340                 345                 350
```

His Gly Ser Val Gly Gln Cys Trp Cys Val Asp Arg Tyr Gly Asn Glu
355 360 365
Val Met Gly Ser Arg Ile Asn Gly Val Ala Asp Cys Ala Ile Asp Phe
370 375 380
Glu Ile Ser Gly Asp Phe Ala Ser Gly Asp Phe His Glu Trp Thr Asp
385 390 395 400
Asp Glu Asp Asp Glu Asp Asp Ile Met Asn Asp Glu Asp Glu Ile Glu
405 410 415
Asp Asp Asp Glu Asp Glu Gly Asp Asp Asp Asp Gly Gly Asp Asp His
420 425 430
Asp Val Tyr Ile *
435 436

<210> 1432
<211> 53
<212> PRT
<213> Homo sapiens

<400> 1432
Met Ser Tyr Val Glu Ile Leu Ile Pro Val Leu Leu Cys Leu His Ala
1 5 10 15
Phe Phe Pro Ser Ser Arg Arg His Val Ala Trp Phe Leu Ile Phe Ile
20 25 30
Cys Lys Phe Phe Lys Phe Cys Leu Ile Leu Lys Phe Ile Ile Leu Ile
35 40 45
Leu Asn Tyr Leu *
50 52

<210> 1433
<211> 76
<212> PRT
<213> Homo sapiens

<400> 1433
Met Glu Leu Lys Gly Phe Trp Leu Cys Leu Phe Leu Arg Phe Val Lys
1 5 10 15
Trp Phe Val Asn Lys Gly Met Ile Leu Cys Thr Leu Phe Tyr Asn Leu
20 25 30
Ile Tyr Ser Leu Tyr Asn Met Cys Trp Thr Val Leu Trp Ile Arg Lys
35 40 45
Tyr Gln Thr Leu Leu Lys Glu Ser Phe Phe Ser Leu Asn Thr Phe Leu
50 55 60
Phe Lys Asp Lys Ala Ser Thr Ser Ile Pro Leu *
65 70 75

<210> 1434
<211> 169
<212> PRT
<213> Homo sapiens

<400> 1434
Met Glu Ser Trp Trp Gly Leu Pro Cys Leu Ala Phe Leu Cys Phe Leu
1 5 10 15
Met His Ala Arg Gly Gln Arg Asp Phe Asp Leu Ala Asp Ala Leu Asp
20 25 30
Asp Pro Glu Pro Thr Lys Lys Pro Asn Ser Asp Ile Tyr Pro Lys Pro
35 40 45
Lys Pro Pro Tyr Tyr Pro Gln Pro Glu Asn Pro Asp Ser Gly Gly Asn
50 55 60
Ile Tyr Pro Arg Pro Lys Pro Arg Pro Gln Pro Gln Pro Gly Asn Ser
65 70 75 80
Gly Asn Ser Gly Gly Ser Tyr Phe Asn Asp Val Asp Arg Asp Asp Gly
85 90 95
Arg Tyr Pro Pro Arg Pro Arg Pro Arg Pro Pro Ala Gly Gly Gly Gly
100 105 110
Gly Gly Tyr Ser Ser Tyr Gly Asn Ser Asp Asn Thr His Gly Gly Asp
115 120 125
His His Ser Thr Tyr Gly Asn Pro Glu Gly Asn Met Val Ala Lys Ile
130 135 140
Val Ser Pro Ile Val Ser Val Val Val Val Thr Leu Leu Gly Ala Ala
145 150 155 160
Ala Gln Leu Phe Gln Thr Lys Gln *
165 168

<210> 1435
<211> 162
<212> PRT
<213> Homo sapiens

<400> 1435
Met Arg Phe Val Thr Leu Ser Ser Ala Cys Leu Cys Pro Cys Pro Leu
1 5 10 15
Gly Pro Cys Trp Thr Arg His Pro Ser Tyr Gly Asn Leu His Glu Ala
20 25 30
Ser Thr Ser Leu Pro Pro Arg His Trp Thr Gly Ala Arg Lys Trp Asn
35 40 45
Glu Ser Ser His Cys Leu Lys Ser Trp Arg Pro Ser Ser Ala Ser Gly
50 55 60
Ser Pro Glu Asn Leu Gly Ser Asp Arg Arg Thr Glu Thr Glu Gly Arg
65 70 75 80
Glu Arg Asp Cys Asp Arg Glu Ala Glu Glu Gly Asp Arg Val Arg Glu
85 90 95
Glu Gln Asn Ser Leu Gln Trp Glu Gln Arg Gln Lys Cys Gly Gly Pro
100 105 110
Thr Gly Arg Gly Gly Arg Glu Gly Glu Gly Arg Arg Glu Gly Gln Leu
115 120 125
Pro Val Gln Val Ala Val Arg Ala Leu Gly Leu Gly Arg Gly Thr Leu
130 135 140
Leu Leu Leu Ala Ser His Thr Gly Ser Ile Arg Gly Pro Arg Glu Gln
145 150 155 160
Val Ser
162

<210> 1436

<211> 77
<212> PRT
<213> Homo sapiens

<400> 1436
Met Trp Ile Val Leu Leu Gly Gly Phe Val Gly Pro Leu Tyr Leu Thr
1 5 10 15
Pro Ala Pro Ser Pro Cys Thr His Thr Leu Gly Val Arg Ala Val Pro
20 25 30
Leu Val Thr Gly Leu Thr Ser Gln Leu Trp Leu Asn Ala Ala Gly Glu
35 40 45
Ser Leu Thr Tyr Arg Met Trp Ser Met Ala Ser Met Thr Glu Gln Pro
50 55 60
Glu Leu Ser Glu Met Tyr Met Leu Pro Thr Leu His Glu
65 70 75 77

<210> 1437
<211> 85
<212> PRT
<213> Homo sapiens

<400> 1437
Met Cys Ser Leu Pro Arg His Leu Leu Phe Leu Ile Ile Phe Arg Ala
1 5 10 15
Tyr Ser Leu Ala Val Asp Leu Ser Thr His Ser Leu Thr Thr Ala Lys
20 25 30
Phe Pro Ser Pro Ile Val Leu Pro Thr Leu Tyr Arg Ser Val Ile Val
35 40 45
Ala Gly Ile Trp Lys Pro Ser Ser Asp Thr Ser Ser Pro Gly Pro Ser
50 55 60
Phe Ser Ser Ile Glu Leu Gln Thr Leu Val Asp Ala Ser Asp Val Glu
65 70 75 80
Glu Pro Pro Cys *
84

<210> 1438
<211> 76
<212> PRT
<213> Homo sapiens

<400> 1438
Met Ile Gly Asp Ile Leu Leu Phe Gly Thr Leu Leu Met Asn Ala Gly
1 5 10 15
Ala Val Leu Asn Phe Lys Leu Lys Lys Lys Asp Thr Gln Gly Phe Gly
20 25 30
Glu Glu Ser Arg Glu Pro Ser Thr Gly Asp Asn Ile Arg Glu Phe Leu
35 40 45
Leu Ser Leu Arg Tyr Phe Arg Ile Phe Ile Ala Leu Trp Asn Ile Phe
50 55 60
Met Met Phe Cys Met Ile Val Leu Phe Gly Ser *
65 70 75

<210> 1439
<211> 425
<212> PRT
<213> Homo sapiens

<400> 1439
Met Ser Leu Thr Ile Trp Thr Val Cys Gly Val Leu Ser Leu Phe Gly
1 5 10 15
Ala Leu Ser Tyr Ala Glu Leu Gly Thr Thr Ile Lys Lys Ser Gly Gly
20 25 30
His Tyr Thr Tyr Ile Leu Glu Val Phe Gly Pro Leu Pro Ala Phe Val
35 40 45
Arg Val Trp Val Glu Leu Leu Ile Ile Arg Pro Ala Ala Thr Ala Val
50 55 60
Ile Ser Leu Ala Phe Gly Arg Tyr Ile Leu Glu Pro Phe Phe Ile Gln
65 70 75 80
Cys Glu Ile Pro Glu Leu Ala Ile Lys Leu Ile Thr Ala Val Gly Ile
85 90 95
Thr Val Val Met Val Leu Asn Ser Met Ser Val Ser Trp Ser Ala Arg
100 105 110
Ile Gln Ile Phe Leu Thr Phe Cys Lys Leu Thr Ala Ile Leu Ile Ile
115 120 125
Ile Val Pro Gly Val Met Gln Leu Ile Lys Gly Gln Thr Gln Asn Phe
130 135 140
Lys Asp Ala Phe Ser Gly Arg Asp Ser Ser Ile Thr Arg Leu Pro Leu
145 150 155 160
Ala Phe Tyr Tyr Gly Met Tyr Ala Tyr Ala Gly Trp Phe Tyr Leu Asn
165 170 175
Phe Val Thr Glu Glu Val Glu Asn Pro Glu Lys Thr Ile Pro Leu Ala
180 185 190
Ile Cys Ile Ser Met Ala Ile Val Thr Ile Gly Tyr Val Leu Thr Asn
195 200 205
Val Ala Tyr Phe Thr Thr Ile Asn Ala Glu Glu Leu Leu Leu Ser Asn
210 215 220
Ala Val Ala Val Thr Phe Ser Glu Arg Leu Leu Gly Asn Phe Ser Leu
225 230 235 240
Ala Val Pro Ile Phe Val Ala Leu Ser Cys Phe Gly Ser Met Asn Gly
245 250 255
Gly Val Phe Ala Val Ser Arg Leu Phe Tyr Val Ala Ser Arg Glu Gly
260 265 270
His Leu Pro Glu Ile Leu Ser Met Ile His Val Arg Lys His Thr Pro
275 280 285
Leu Pro Ala Val Ile Val Leu His Pro Leu Thr Met Ile Met Leu Phe
290 295 300
Ser Gly Asp Leu Asp Ser Leu Leu Asn Phe Leu Ser Phe Ala Arg Trp
305 310 315 320
Leu Phe Ile Gly Leu Ala Val Ala Gly Leu Ile Tyr Leu Arg Tyr Lys
325 330 335
Cys Pro Asp Met His Arg Pro Phe Lys Val Pro Leu Phe Ile Pro Ala
340 345 350
Leu Phe Ser Phe Thr Cys Leu Phe Met Val Ala Leu Ser Leu Tyr Ser
355 360 365
Asp Pro Phe Ser Thr Gly Ile Gly Phe Val Ile Thr Leu Thr Gly Val
370 375 380
Pro Ala Tyr Tyr Leu Phe Ile Ile Trp Asp Lys Lys Pro Arg Trp Phe
385 390 395 400

Arg Ile Met Ser Glu Lys Ile Thr Arg Thr Leu Gln Ile Ile Leu Glu
405 410 415
Val Val Pro Glu Glu Asp Lys Leu *
420 424

<210> 1440
<211> 70
<212> PRT
<213> Homo sapiens

<400> 1440
Met Ser Val Phe Trp Gly Phe Val Gly Phe Leu Val Pro Trp Phe Ile
1 5 10 15
Pro Lys Gly Pro Asn Arg Gly Val Ile Ile Thr Met Leu Val Thr Cys
20 25 30
Ser Val Cys Cys Tyr Leu Phe Trp Leu Ile Ala Ile Leu Ala Gln Leu
35 40 45
Asn Pro Leu Phe Gly Pro Gln Leu Lys Asn Glu Thr Ile Trp Tyr Leu
50 55 60
Lys Tyr His Trp Pro *
65 69

<210> 1441
<211> 1691
<212> PRT
<213> Homo sapiens

<400> 1441
Met Trp Ser Leu His Ile Val Leu Met Arg Cys Ser Phe Arg Leu Thr
1 5 10 15
Lys Ser Leu Ala Thr Gly Pro Trp Ser Leu Ile Leu Ile Leu Phe Ser
20 25 30
Val Gln Tyr Val Tyr Gly Ser Gly Lys Lys Tyr Ile Gly Pro Cys Gly
35 40 45
Gly Arg Asp Cys Ser Val Cys His Cys Val Pro Glu Lys Gly Ser Arg
50 55 60
Gly Pro Pro Gly Pro Pro Gly Pro Gln Gly Pro Ile Gly Pro Leu Gly
65 70 75 80
Ala Pro Gly Pro Ile Gly Leu Ser Gly Glu Lys Gly Met Arg Gly Asp
85 90 95
Arg Gly Pro Pro Gly Ala Ala Gly Asp Lys Gly Asp Lys Gly Pro Thr
100 105 110
Gly Val Pro Gly Phe Pro Gly Leu Asp Gly Ile Pro Gly His Pro Gly
115 120 125
Pro Pro Gly Pro Arg Gly Lys Pro Gly Met Ser Gly His Asn Gly Ser
130 135 140
Arg Gly Asp Pro Gly Phe Pro Gly Gly Arg Gly Ala Leu Gly Pro Gly
145 150 155 160
Gly Pro Leu Gly His Pro Gly Glu Lys Gly Glu Lys Gly Asn Ser Val
165 170 175
Phe Ile Leu Gly Ala Val Lys Gly Ile Gln Gly Asp Arg Gly Asp Pro
180 185 190
Gly Leu Pro Gly Leu Pro Gly Ser Trp Gly Ala Gly Gly Pro Ala Gly

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        195                 200                 205
Pro Thr Gly Tyr Pro Gly Glu Pro Gly Leu Val Gly Pro Pro Gly Gln
    210                 215                 220
Pro Gly Arg Pro Gly Leu Lys Gly Asn Pro Gly Val Gly Val Lys Gly
225                 230                 235                 240
Gln Met Gly Asp Pro Gly Glu Val Gly Gln Gln Gly Ser Pro Gly Pro
                245                 250                 255
Thr Leu Leu Val Glu Pro Pro Asp Phe Cys Leu Tyr Lys Gly Glu Lys
            260                 265                 270
Gly Ile Lys Gly Ile Pro Gly Met Val Gly Leu Pro Gly Pro Pro Gly
        275                 280                 285
Arg Lys Gly Glu Ser Gly Ile Gly Ala Lys Gly Glu Lys Gly Ile Pro
    290                 295                 300
Gly Phe Pro Gly Pro Arg Gly Asp Pro Gly Ser Tyr Gly Ser Pro Gly
305                 310                 315                 320
Phe Pro Gly Leu Lys Gly Glu Leu Gly Leu Val Gly Asp Pro Gly Leu
                325                 330                 335
Phe Gly Leu Ile Gly Pro Lys Gly Asp Pro Gly Asn Arg Gly His Pro
            340                 345                 350
Gly Pro Pro Gly Val Leu Val Thr Pro Pro Leu Pro Leu Lys Gly Pro
        355                 360                 365
Pro Gly Asp Pro Gly Phe Pro Gly Arg Tyr Gly Glu Thr Gly Asp Val
    370                 375                 380
Gly Pro Pro Gly Pro Pro Gly Leu Leu Gly Arg Pro Gly Glu Ala Cys
385                 390                 395                 400
Ala Gly Met Ile Gly Pro Pro Gly Pro Gln Gly Phe Pro Gly Leu Pro
                405                 410                 415
Gly Leu Pro Gly Glu Ala Gly Ile Pro Gly Arg Pro Asp Ser Ala Pro
            420                 425                 430
Gly Lys Pro Gly Lys Pro Gly Ser Pro Gly Leu Pro Gly Ala Pro Gly
        435                 440                 445
Leu Gln Gly Leu Pro Gly Ser Ser Val Ile Tyr Cys Ser Val Gly Asn
    450                 455                 460
Pro Gly Pro Gln Gly Ile Lys Gly Lys Val Gly Pro Pro Gly Gly Arg
465                 470                 475                 480
Gly Pro Lys Gly Glu Lys Gly Asn Glu Gly Leu Cys Ala Cys Glu Pro
                485                 490                 495
Gly Pro Met Gly Pro Pro Gly Pro Pro Gly Leu Pro Gly Arg Gln Gly
            500                 505                 510
Ser Lys Gly Asp Leu Gly Leu Pro Gly Trp Leu Gly Thr Lys Gly Asp
        515                 520                 525
Pro Gly Pro Pro Gly Ala Glu Gly Pro Pro Gly Leu Pro Gly Lys His
    530                 535                 540
Gly Ala Ser Gly Pro Pro Gly Asn Lys Gly Ala Lys Gly Asp Met Val
545                 550                 555                 560
Val Ser Arg Val Lys Gly His Lys Gly Glu Arg Gly Pro Asp Gly Pro
                565                 570                 575
Pro Gly Phe Pro Gly Gln Pro Gly Ser His Gly Arg Asp Gly His Ala
            580                 585                 590
Gly Glu Lys Gly Asp Pro Gly Pro Pro Gly Asp His Glu Asp Ala Thr
        595                 600                 605
Pro Gly Gly Lys Gly Phe Pro Gly Pro Leu Gly Pro Pro Gly Lys Ala
    610                 615                 620
Gly Pro Val Gly Pro Pro Gly Leu Gly Phe Pro Gly Pro Pro Gly Glu
625                 630                 635                 640
Arg Gly His Pro Gly Val Pro Gly His Pro Gly Val Arg Gly Pro Asp
                645                 650                 655
Gly Leu Lys Gly Gln Lys Gly Asp Thr Ile Ser Cys Asn Val Thr Tyr
            660                 665                 670
```

Pro Gly Arg His Gly Pro Pro Gly Phe Asp Gly Pro Pro Gly Pro Lys
675 680 685
Gly Phe Pro Gly Pro Gln Gly Ala Pro Gly Leu Ser Gly Ser Asp Gly
690 695 700
His Lys Gly Arg Pro Gly Thr Pro Gly Thr Ala Glu Ile Pro Gly Pro
705 710 715 720
Pro Gly Phe Arg Gly Asp Met Gly Asp Pro Gly Phe Gly Gly Glu Lys
725 730 735
Gly Ser Ser Pro Val Gly Pro Pro Gly Pro Pro Gly Ser Pro Gly Val
740 745 750
Asn Gly Gln Lys Gly Ile Pro Gly Asp Pro Ala Phe Gly His Leu Gly
755 760 765
Pro Pro Gly Lys Arg Gly Leu Ser Gly Val Pro Gly Ile Lys Gly Pro
770 775 780
Arg Gly Asp Pro Gly Cys Pro Gly Ala Glu Gly Pro Ala Gly Ile Pro
785 790 795 800
Gly Phe Leu Gly Leu Lys Gly Pro Lys Gly Arg Glu Gly His Ala Gly
805 810 815
Phe Pro Gly Val Pro Gly Pro Pro Gly His Ser Cys Glu Arg Gly Ala
820 825 830
Pro Gly Ile Pro Gly Gln Pro Gly Leu Pro Gly Tyr Pro Gly Ser Pro
835 840 845
Gly Ala Pro Gly Gly Lys Gly Gln Pro Gly Asp Val Gly Pro Pro Gly
850 855 860
Pro Ala Gly Met Lys Gly Leu Pro Gly Leu Pro Gly Arg Pro Gly Ala
865 870 875 880
His Gly Pro Pro Gly Leu Pro Gly Ile Pro Gly Pro Phe Gly Asp Asp
885 890 895
Gly Leu Pro Gly Pro Pro Gly Pro Lys Gly Pro Arg Gly Leu Pro Gly
900 905 910
Phe Pro Gly Phe Pro Gly Glu Arg Gly Lys Pro Gly Ala Glu Gly Cys
915 920 925
Pro Gly Ala Lys Gly Glu Pro Gly Glu Lys Gly Met Ser Gly Leu Pro
930 935 940
Gly Asp Arg Gly Leu Arg Gly Ala Lys Gly Ala Ile Gly Pro Pro Gly
945 950 955 960
Asp Glu Gly Glu Met Ala Ile Ile Ser Gln Lys Gly Thr Pro Gly Glu
965 970 975
Pro Gly Pro Pro Gly Asp Asp Gly Phe Pro Gly Glu Arg Gly Asp Lys
980 985 990
Gly Thr Pro Gly Met Gln Gly Arg Arg Gly Glu Leu Gly Arg Tyr Gly
995 1000 1005
Pro Pro Gly Phe His Arg Gly Glu Pro Gly Glu Lys Gly Gln Pro Gly
1010 1015 1020
Pro Pro Gly Pro Pro Gly Pro Pro Gly Ser Thr Gly Leu Arg Gly Phe
1025 1030 1035 1040
Ile Gly Phe Pro Gly Leu Pro Gly Asp Gln Gly Glu Pro Gly Ser Pro
1045 1050 1055
Gly Pro Pro Gly Phe Ser Gly Ile Asp Gly Ala Arg Gly Pro Lys Gly
1060 1065 1070
Asn Lys Gly Asp Pro Ala Ser His Phe Gly Pro Pro Gly Pro Lys Gly
1075 1080 1085
Glu Pro Gly Ser Pro Gly Cys Pro Gly His Phe Gly Ala Ser Gly Glu
1090 1095 1100
Gln Gly Leu Pro Gly Ile Gln Gly Pro Arg Gly Ser Pro Gly Arg Pro
1105 1110 1115 1120
Gly Pro Pro Gly Ser Ser Gly Pro Pro Gly Cys Pro Gly Asp His Gly
1125 1130 1135
Met Pro Gly Leu Arg Gly Gln Pro Gly Glu Met Gly Asp Pro Gly Pro

1140 1145 1150
Arg Gly Leu Gln Gly Asp Pro Gly Ile Pro Gly Pro Pro Gly Ile Lys
1155 1160 1165
Gly Pro Ser Gly Ser Pro Gly Leu Asn Gly Leu His Gly Leu Lys Gly
1170 1175 1180
Gln Lys Gly Thr Lys Gly Ala Ser Gly Leu His Asp Val Gly Pro Pro
1185 1190 1195 1200
Gly Pro Val Gly Ile Pro Gly Leu Lys Gly Glu Arg Gly Asp Pro Gly
1205 1210 1215
Ser Pro Gly Ile Ser Pro Pro Gly Pro Arg Gly Lys Lys Gly Pro Pro
1220 1225 1230
Gly Pro Pro Gly Ser Ser Gly Pro Pro Gly Pro Ala Gly Ala Thr Gly
1235 1240 1245
Arg Ala Pro Lys Asp Ile Pro Asp Pro Gly Pro Pro Gly Asp Gln Gly
1250 1255 1260
Pro Pro Gly Pro Asp Gly Pro Arg Gly Ala Pro Gly Pro Pro Gly Leu
1265 1270 1275 1280
Pro Gly Ser Val Asp Leu Leu Arg Gly Glu Pro Gly Asp Cys Gly Leu
1285 1290 1295
Pro Gly Pro Pro Gly Pro Pro Gly Pro Pro Gly Pro Pro Gly Tyr Lys
1300 1305 1310
Gly Phe Pro Gly Cys Asp Gly Lys Asp Gly Gln Lys Gly Pro Val Gly
1315 1320 1325
Phe Pro Gly Pro Gln Gly Pro His Gly Phe Pro Gly Pro Pro Gly Glu
1330 1335 1340
Lys Gly Leu Pro Gly Pro Pro Gly Arg Lys Gly Pro Thr Gly Leu Pro
1345 1350 1355 1360
Gly Pro Arg Gly Glu Pro Gly Pro Pro Ala Asp Val Asp Asp Cys Pro
1365 1370 1375
Arg Ile Pro Gly Leu Pro Gly Ala Pro Gly Met Arg Gly Pro Glu Gly
1380 1385 1390
Ala Met Gly Leu Pro Gly Met Arg Gly Pro Ser Gly Pro Gly Cys Lys
1395 1400 1405
Gly Glu Pro Gly Leu Asp Gly Arg Arg Gly Val Asp Gly Val Pro Gly
1410 1415 1420
Ser Pro Gly Pro Pro Gly Arg Lys Gly Asp Thr Gly Glu Asp Gly Tyr
1425 1430 1435 1440
Pro Gly Gly Pro Gly Pro Pro Gly Pro Ile Gly Asp Pro Gly Pro Lys
1445 1450 1455
Gly Phe Gly Pro Gly Tyr Leu Gly Gly Phe Leu Leu Val Leu His Ser
1460 1465 1470
Gln Thr Asp Gln Glu Pro Thr Cys Pro Leu Gly Met Pro Arg Leu Trp
1475 1480 1485
Thr Gly Tyr Ser Leu Leu Tyr Leu Glu Gly Gln Glu Lys Ala His Asn
1490 1495 1500
Gln Asp Leu Gly Leu Ala Gly Ser Cys Leu Pro Val Phe Ser Thr Leu
1505 1510 1515 1520
Pro Phe Ala Tyr Cys Asn Ile His Gln Val Cys His Tyr Ala Gln Arg
1525 1530 1535
Asn Asp Arg Ser Tyr Trp Leu Ala Ser Ala Ala Pro Leu Pro Met Met
1540 1545 1550
Pro Leu Ser Glu Glu Ala Ile Arg Pro Tyr Val Ser Arg Cys Ala Val
1555 1560 1565
Cys Glu Ala Pro Ala Gln Ala Val Ala Val His Ser Gln Asp Gln Ser
1570 1575 1580
Ile Pro Pro Cys Pro Gln Thr Trp Arg Ser Leu Trp Ile Gly Tyr Ser
1585 1590 1595 1600
Phe Leu Met His Thr Gly Ala Gly Asp Gln Gly Gly Gly Gln Ala Leu
1605 1610 1615

Met Ser Pro Gly Ser Cys Leu Glu Asp Phe Arg Ala Ala Pro Phe Leu
1620 1625 1630
Glu Cys Gln Gly Arg Gln Gly Thr Cys His Phe Phe Ala Asn Lys Tyr
1635 1640 1645
Ser Phe Trp Leu Thr Thr Val Lys Ala Asp Phe Glu Phe Ser Ser Ala
1650 1655 1660
Pro Ala Pro Asp Thr Leu Lys Glu Ser Gln Ala Gln Arg Gln Lys Ile
1665 1670 1675 1680
Ser Arg Cys Gln Val Cys Val Lys Tyr Ser *
1685 1690

<210> 1442
<211> 153
<212> PRT
<213> Homo sapiens

<400> 1442
Met Gly Val Met Ala Pro Arg Thr Leu Leu Leu Leu Leu Leu Gly Ala
1 5 10 15
Leu Ala Leu Thr Glu Thr Trp Ala Gly Glu Cys Gly Val Gly Arg Glu
20 25 30
Arg Ala Ser Ala Gly Arg Ser Glu Trp Pro Ala Arg Pro Gly Glu Pro
35 40 45
Arg Arg Glu Glu Gly Arg Ala Gly Leu Ser Leu Ser Ser Pro Pro Gly
50 55 60
Ser His Ser Leu Arg Tyr Phe Ser Thr Ala Val Ser Gln Pro Gly Arg
65 70 75 80
Gly Glu Pro Arg Phe Ile Ala Val Gly Tyr Val Asp Asp Thr Glu Phe
85 90 95
Val Arg Phe Asp Ser Asp Ser Val Ser Pro Arg Met Glu Arg Arg Ala
100 105 110
Pro Trp Val Glu Gln Glu Gly Leu Glu Tyr Trp Asp Gln Glu Thr Arg
115 120 125
Asn Ala Lys Gly His Ala Gln Ile Tyr Arg Val Asn Leu Arg Thr Leu
130 135 140
Leu Arg Tyr Tyr Asn Gln Ser Glu Ala
145 150 153

<210> 1443
<211> 58
<212> PRT
<213> Homo sapiens

<400> 1443
Met Ser Leu Leu Cys Leu Lys Phe Phe Ser Gly Leu Trp Thr Ile Thr
1 5 10 15
Phe Ser Lys Gly Ala Lys Ile Ile His Trp Gly Arg Ser Leu Phe Asn
20 25 30
Trp Ile Ser Met Cys Lys Arg Met Lys Leu Asp Pro Tyr Ser Tyr His
35 40 45
Thr Gln Lys Leu Thr Gln Asn Gly Ser *
50 55 57

<210> 1444
<211> 69
<212> PRT
<213> Homo sapiens

<400> 1444
Met Pro Val Pro Leu Ala Tyr Phe Gln Ser Ser Ile Val Leu Phe Pro
1 5 10 15
Leu Ile Phe Ser Leu Val Thr Cys Val Ser Leu Asp Gly Glu Pro Lys
20 25 30
Ser Val Val Gly Val Ile Ser Ile Ser Ala Tyr Tyr Arg Ala Ile Ser
35 40 45
Ile Leu Leu Ile Phe Ser Lys Ser Phe Cys Cys Ala Ser Leu Ala Gly
50 55 60
Val Leu Val Ile *
65 68

<210> 1445
<211> 826
<212> PRT
<213> Homo sapiens

<400> 1445
Met Gly Trp Leu Cys Ser Gly Leu Leu Phe Pro Val Ser Cys Leu Val
1 5 10 15
Leu Leu Gln Val Ala Ser Ser Gly Asn Met Lys Val Leu Gln Glu Pro
20 25 30
Thr Cys Val Ser Asp Tyr Met Ser Ile Ser Thr Cys Glu Trp Lys Met
35 40 45
Asn Gly Pro Thr Asn Cys Ser Thr Glu Leu Arg Leu Leu Tyr Gln Leu
50 55 60
Val Phe Leu Leu Ser Glu Ala His Thr Cys Val Pro Glu Asn Asn Gly
65 70 75 80
Gly Ala Gly Cys Val Cys His Leu Leu Met Asp Asp Val Val Ser Ala
85 90 95
Asp Asn Tyr Thr Leu Asp Leu Trp Ala Gly Gln Gln Leu Leu Trp Lys
100 105 110
Gly Ser Phe Lys Pro Ser Glu His Val Lys Pro Arg Ala Pro Gly Asn
115 120 125
Leu Thr Val His Thr Asn Val Ser Asp Thr Leu Leu Leu Thr Trp Ser
130 135 140
Asn Pro Tyr Pro Pro Asp Asn Tyr Leu Tyr Asn His Leu Thr Tyr Ala
145 150 155 160
Val Asn Ile Trp Ser Glu Asn Asp Pro Ala Asp Phe Arg Ile Tyr Asn
165 170 175
Val Thr Tyr Leu Glu Pro Ser Leu Arg Ile Ala Ala Ser Thr Leu Lys
180 185 190
Ser Gly Ile Ser Tyr Arg Ala Arg Val Arg Ala Trp Ala Gln Cys Tyr
195 200 205
Asn Thr Thr Trp Ser Glu Trp Ser Pro Ser Thr Lys Trp His Asn Ser
210 215 220
Tyr Arg Glu Pro Phe Glu Gln His Leu Leu Leu Gly Val Ser Val Ser
225 230 235 240

Cys Ile Val Ile Leu Ala Val Cys Leu Leu Cys Tyr Val Ser Ile Thr
245 250 255
Lys Ile Lys Lys Glu Trp Trp Asp Gln Ile Pro Asn Pro Ala Arg Ser
260 265 270
Arg Leu Val Ala Ile Ile Ile Gln Asp Ala Gln Gly Ser Gln Trp Glu
275 280 285
Lys Arg Ser Arg Gly Gln Glu Pro Ala Lys Cys Pro His Trp Lys Asn
290 295 300
Cys Leu Thr Lys Leu Leu Pro Cys Phe Leu Glu His Asn Met Lys Arg
305 310 315 320
Asp Glu Asp Pro His Lys Ala Ala Lys Glu Met Pro Phe Gln Gly Ser
325 330 335
Gly Lys Ser Ala Trp Cys Pro Val Glu Ile Ser Lys Thr Val Leu Trp
340 345 350
Pro Glu Ser Ile Ser Val Val Arg Cys Val Glu Leu Phe Glu Ala Pro
355 360 365
Val Glu Cys Glu Glu Glu Glu Glu Val Glu Glu Glu Lys Gly Ser Phe
370 375 380
Cys Ala Ser Pro Glu Ser Ser Arg Asp Asp Phe Gln Glu Gly Arg Glu
385 390 395 400
Gly Ile Val Ala Arg Leu Thr Glu Ser Leu Phe Leu Asp Leu Leu Gly
405 410 415
Glu Glu Asn Gly Gly Phe Cys Gln Gln Asp Met Gly Glu Ser Cys Leu
420 425 430
Leu Pro Pro Ser Gly Ser Thr Ser Ala His Met Pro Trp Asp Glu Phe
435 440 445
Pro Ser Ala Gly Pro Lys Glu Ala Pro Pro Trp Gly Lys Glu Gln Pro
450 455 460
Leu His Leu Glu Pro Ser Pro Pro Ala Ser Pro Thr Gln Ser Pro Asp
465 470 475 480
Asn Leu Thr Cys Thr Glu Thr Pro Leu Val Ile Ala Gly Asn Pro Ala
485 490 495
Tyr Arg Ser Phe Ser Asn Ser Leu Ser Gln Ser Pro Cys Pro Arg Glu
500 505 510
Leu Gly Pro Asp Pro Leu Leu Ala Arg His Leu Glu Glu Val Glu Pro
515 520 525
Glu Met Pro Cys Val Pro Gln Leu Ser Glu Pro Thr Thr Val Pro Gln
530 535 540
Pro Glu Pro Glu Thr Trp Glu Gln Ile Leu Arg Arg Asn Val Leu Gln
545 550 555 560
His Gly Ala Ala Ala Ala Pro Val Ser Ala Pro Thr Ser Gly Tyr Gln
565 570 575
Glu Phe Val His Ala Val Glu Gln Gly Gly Thr Gln Ala Ser Ala Val
580 585 590
Val Gly Leu Gly Pro Pro Gly Glu Ala Gly Tyr Lys Ala Phe Ser Ser
595 600 605
Leu Leu Ala Ser Ser Ala Val Ser Pro Glu Lys Cys Gly Phe Gly Ala
610 615 620
Ser Ser Gly Glu Glu Gly Tyr Lys Pro Phe Gln Asp Leu Ile Pro Gly
625 630 635 640
Cys Pro Gly Asp Pro Ala Pro Val Pro Val Pro Leu Phe Thr Phe Gly
645 650 655
Leu Asp Arg Glu Pro Pro Arg Ser Pro Gln Ser Ser His Leu Pro Ser
660 665 670
Ser Ser Pro Glu His Leu Gly Leu Glu Pro Gly Glu Lys Val Glu Asp
675 680 685
Met Pro Lys Pro Pro Leu Pro Gln Glu Gln Ala Thr Asp Pro Leu Val
690 695 700
Asp Ser Leu Gly Ser Gly Ile Val Tyr Ser Ala Leu Thr Cys His Leu

705 710 715 720
Cys Gly His Leu Lys Gln Cys His Gly Gln Glu Asp Gly Gly Gln Thr
725 730 735
Pro Val Met Ala Ser Pro Cys Cys Gly Cys Cys Cys Gly Asp Arg Ala
740 745 750
Ser Pro Pro Thr Thr Pro Leu Arg Ala Pro Asp Pro Ser Pro Gly Gly
755 760 765
Val Pro Leu Glu Ala Ser Leu Cys Pro Ala Ser Leu Ala Pro Ser Gly
770 775 780
Ile Ser Glu Lys Ser Lys Ser Ser Ser Ser Phe His Pro Ala Pro Gly
785 790 795 800
Asn Ala Gln Ser Ser Ser Gln Thr Pro Lys Ile Val Asn Phe Val Ser
805 810 815
Val Gly Pro Thr Tyr Met Arg Val Ser *
820 825

<210> 1446
<211> 367
<212> PRT
<213> Homo sapiens

<400> 1446
Met Ala Leu Arg Phe Leu Leu Gly Phe Leu Leu Ala Gly Val Asp Leu
1 5 10 15
Gly Val Tyr Leu Met Arg Leu Glu Leu Cys Asp Pro Thr Gln Arg Leu
20 25 30
Arg Val Ala Leu Ala Gly Glu Leu Val Gly Val Gly Gly His Phe Leu
35 40 45
Phe Leu Gly Leu Ala Leu Val Ser Lys Asp Trp Arg Phe Leu Gln Arg
50 55 60
Met Ile Thr Ala Pro Cys Ile Leu Phe Leu Phe Tyr Gly Trp Pro Gly
65 70 75 80
Leu Phe Leu Glu Ser Ala Arg Trp Leu Ile Val Lys Arg Gln Ile Glu
85 90 95
Glu Ala Gln Ser Val Leu Arg Ile Leu Ala Glu Arg Asn Arg Pro His
100 105 110
Gly Gln Met Leu Gly Glu Glu Ala Gln Glu Ala Leu Gln Asp Leu Glu
115 120 125
Asn Thr Cys Pro Leu Pro Ala Thr Ser Ser Phe Ser Phe Ala Ser Leu
130 135 140
Leu Asn Tyr Arg Asn Ile Trp Lys Asn Leu Leu Ile Leu Gly Phe Thr
145 150 155 160
Asn Phe Ile Ala His Ala Ile Arg His Cys Tyr Gln Pro Val Gly Gly
165 170 175
Gly Gly Ser Pro Ser Asp Phe Tyr Leu Cys Ser Leu Leu Ala Ser Gly
180 185 190
Thr Ala Ala Leu Ala Cys Val Phe Leu Gly Val Thr Val Asp Arg Phe
195 200 205
Gly Arg Arg Gly Ile Leu Leu Leu Ser Met Thr Leu Thr Gly Ile Ala
210 215 220
Ser Leu Val Leu Leu Gly Leu Trp Asp Tyr Leu Asn Glu Ala Ala Ile
225 230 235 240
Thr Thr Phe Ser Val Leu Gly Leu Phe Ser Ser Gln Ala Ala Ala Ile
245 250 255
Leu Ser Thr Leu Leu Ala Ala Glu Val Ile Pro Thr Thr Val Arg Gly
260 265 270

Arg Gly Leu Gly Leu Ile Met Ala Leu Gly Ala Leu Gly Gly Leu Ser
275 280 285
Gly Pro Ala Gln Arg Leu His Met Gly His Gly Ala Phe Leu Gln His
290 295 300
Val Val Leu Ala Ala Cys Ala Leu Leu Cys Ile Leu Ser Ile Met Leu
305 310 315 320
Leu Pro Glu Thr Lys Arg Lys Leu Leu Pro Glu Val Leu Arg Asp Gly
325 330 335
Glu Leu Cys Arg Arg Pro Ser Leu Leu Arg Gln Pro Pro Pro Thr Arg
340 345 350
Cys Asp His Val Pro Leu Leu Ala Thr Pro Asn Pro Ala Leu *
355 360 365 366

<210> 1447
<211> 79
<212> PRT
<213> Homo sapiens

<400> 1447
Met Ala Ile Ser Trp Leu Gly Thr Trp Leu Leu Gln Ser His Arg His
1 5 10 15
Trp Ser Glu Pro Gln Leu Cys Arg Leu Pro Ala Arg His His Leu Ile
20 25 30
Asn Leu Asn Phe Met Val Ala Glu Gly Ile Gly Asp Arg Ala Trp His
35 40 45
Ile Ile Ser Ala Gln Leu Phe Met Thr Phe Ser Phe His Ala Val Ile
50 55 60
Leu Gln Thr Asp Leu Gly Glu Ala Gly Lys Tyr Lys Asp Lys *
65 70 75 78

<210> 1448
<211> 276
<212> PRT
<213> Homo sapiens

<400> 1448
Met Val Trp Val Val Leu Leu Ser Leu Leu Cys Tyr Leu Val Leu Phe
1 5 10 15
Leu Cys Arg His Ser Ser His Arg Gly Val Phe Leu Ser Val Thr Ile
20 25 30
Leu Ile Tyr Leu Leu Met Gly Glu Met His Met Val Asp Thr Val Thr
35 40 45
Trp His Lys Met Arg Gly Ala Gln Met Ile Val Ala Met Lys Ala Val
50 55 60
Ser Leu Gly Phe Asp Leu Asp Arg Gly Glu Val Gly Thr Val Pro Ser
65 70 75 80
Pro Val Glu Phe Met Gly Tyr Leu Tyr Phe Val Gly Thr Ile Val Phe
85 90 95
Gly Pro Trp Ile Ser Phe His Ser Tyr Leu Gln Ala Val Gln Gly Arg
100 105 110
Pro Leu Ser Cys Arg Trp Leu Gln Lys Val Ala Arg Ser Leu Ala Leu
115 120 125
Ala Leu Leu Cys Leu Val Leu Ser Thr Cys Val Gly Pro Tyr Leu Phe

```
    130                 135                 140
Pro Tyr Phe Ile Pro Leu Asn Gly Asp Arg Leu Leu Arg Lys Trp Leu
145                 150                 155                 160
Arg Ala Tyr Glu Ser Ala Val Ser Phe His Phe Ser Asn Tyr Phe Val
                165                 170                 175
Gly Phe Leu Ser Glu Ala Thr Ala Thr Leu Ala Gly Ala Gly Phe Thr
            180                 185                 190
Glu Glu Lys Asp His Leu Glu Trp Asp Leu Thr Val Ser Lys Pro Leu
        195                 200                 205
Asn Val Glu Leu Pro Arg Ser Met Val Glu Val Val Thr Ser Trp Asn
    210                 215                 220
Leu Pro Met Ser Tyr Trp Leu Asn Asn Tyr Gly Phe Lys Asn Ala Leu
225                 230                 235                 240
Arg Leu Gly Thr Leu Leu Gly Cys Ala Gly His Leu Cys Ser Gln Arg
                245                 250                 255
Pro Ser Lys Leu Leu Lys Phe Pro Pro Gly Trp Gly Pro Cys Cys Pro
            260                 265                 270
Gly Phe Leu  *
        275
```

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     <210> 1449
     <211> 597
     <212> PRT
     <213> Homo sapiens


     <400> 1449
Met Glu Phe Gly Leu Ser Trp Val Phe Leu Val Ala Ile Leu Lys Gly
 1               5                  10                  15
Val Gln Cys Glu Val Gln Leu Val Glu Ser Gly Gly Gly Leu Val Gln
            20                  25                  30
Pro Gly Gly Ser Leu Arg Leu Ser Cys Ala Ala Ser Gly Phe Thr Phe
        35                  40                  45
Ser Ser Tyr Trp Met His Trp Val Arg Gln Ala Pro Gly Lys Gly Leu
    50                  55                  60
Val Trp Val Ser Arg Ile Asn Thr Asp Gly Ser Ser Thr Ser Tyr Ala
65                  70                  75                  80
Asp Ser Val Lys Gly Arg Phe Thr Ile Ser Arg Asp Asn Ala Lys Asn
                85                  90                  95
Thr Leu Tyr Leu Gln Met Asn Ser Leu Arg Ala Glu Asp Thr Ala Val
            100                 105                 110
Tyr Tyr Cys Ala Arg Ala Asp Asn Cys Ser Ser Thr Ser Cys Tyr Lys
        115                 120                 125
Cys Phe Asp Tyr Trp Gly Gln Gly Thr Leu Val Thr Val Ser Ser Gly
    130                 135                 140
Ser Ala Ser Ala Pro Thr Leu Phe Pro Leu Val Ser Cys Glu Asn Ser
145                 150                 155                 160
Pro Ser Asp Thr Ser Ser Val Ala Val Gly Cys Leu Ala Gln Asp Phe
                165                 170                 175
Leu Pro Asp Ser Ile Thr Phe Ser Trp Lys Tyr Lys Asn Asn Ser Asp
            180                 185                 190
Ile Ser Ser Thr Arg Gly Phe Pro Ser Val Leu Arg Gly Gly Lys Tyr
        195                 200                 205
Ala Ala Thr Ser Gln Val Leu Leu Pro Ser Lys Asp Val Met Gln Gly
    210                 215                 220
Thr Asp Glu His Val Val Cys Lys Val Gln His Pro Asn Gly Asn Lys
225                 230                 235                 240
```

Glu Lys Asn Val Pro Leu Pro Val Ile Ala Glu Leu Pro Pro Lys Val
245 250 255
Ser Val Phe Val Pro Pro Arg Asp Gly Phe Phe Gly Asn Pro Arg Lys
260 265 270
Ser Lys Leu Ile Cys Gln Ala Thr Gly Phe Ser Pro Arg Gln Ile Gln
275 280 285
Val Ser Trp Leu Arg Glu Gly Lys Gln Val Gly Ser Gly Val Thr Thr
290 295 300
Asp Gln Val Gln Ala Glu Ala Lys Glu Ser Gly Pro Thr Thr Tyr Lys
305 310 315 320
Val Thr Ser Thr Leu Thr Ile Lys Glu Ser Asp Trp Leu Ser Gln Ser
325 330 335
Met Phe Thr Cys Arg Val Asp His Arg Gly Leu Thr Phe Gln Gln Asn
340 345 350
Ala Ser Ser Met Cys Val Pro Asp Gln Asp Thr Ala Ile Arg Val Phe
355 360 365
Ala Ile Pro Pro Ser Phe Ala Ser Ile Phe Leu Thr Lys Ser Thr Lys
370 375 380
Leu Thr Cys Leu Val Thr Asp Leu Thr Thr Tyr Asp Ser Val Thr Ile
385 390 395 400
Ser Trp Thr Arg Gln Asn Gly Glu Ala Val Lys Thr His Thr Asn Ile
405 410 415
Ser Glu Ser His Pro Asn Ala Thr Phe Ser Ala Val Gly Glu Ala Ser
420 425 430
Ile Cys Glu Asp Asp Trp Asn Ser Gly Glu Arg Phe Thr Cys Thr Val
435 440 445
Thr His Thr Asp Leu Pro Ser Pro Leu Lys Gln Thr Ile Ser Arg Pro
450 455 460
Lys Gly Val Ala Leu His Arg Pro Asp Val Tyr Leu Leu Pro Pro Ala
465 470 475 480
Arg Glu Gln Leu Asn Leu Arg Glu Ser Ala Thr Ile Thr Cys Leu Val
485 490 495
Thr Gly Phe Ser Pro Ala Asp Val Phe Val Gln Trp Met Gln Arg Gly
500 505 510
Gln Pro Leu Ser Pro Glu Lys Tyr Val Thr Ser Ala Pro Met Pro Glu
515 520 525
Pro Gln Ala Pro Gly Arg Tyr Phe Ala His Ser Ile Leu Thr Val Ser
530 535 540
Glu Glu Glu Trp Asn Thr Gly Glu Thr Tyr Thr Cys Val Val Ala His
545 550 555 560
Glu Ala Leu Pro Asn Arg Val Thr Glu Arg Thr Val Asp Lys Ser Thr
565 570 575
Gly Lys Pro Thr Leu Tyr Asn Val Ser Leu Val Met Ser Asp Thr Ala
580 585 590
Gly Thr Cys Tyr *
595 596

<210> 1450
<211> 276
<212> PRT
<213> Homo sapiens

<400> 1450
Met Pro Ala Leu Arg Pro Ala Leu Leu Trp Ala Leu Leu Ala Leu Trp
1 5 10 15
Leu Cys Cys Ala Thr Pro Ala His Ala Leu Gln Cys Arg Asp Gly Tyr

```
                20                  25                  30
Glu Pro Cys Val Asn Glu Gly Met Cys Val Thr Tyr His Asn Gly Thr
            35                  40                  45
Gly Tyr Cys Lys Cys Pro Glu Gly Phe Leu Gly Glu Tyr Cys Gln His
    50                  55                  60
Arg Asp Pro Cys Glu Lys Asn Arg Cys Gln Asn Gly Gly Thr Cys Val
65                  70                  75                  80
Ala Gln Ala Met Leu Gly Lys Ala Thr Cys Arg Cys Ala Ser Gly Phe
                85                  90                  95
Thr Gly Glu Asp Cys Gln Tyr Ser Thr Ser His Pro Cys Phe Val Ser
            100                 105                 110
Arg Pro Cys Leu Asn Gly Gly Thr Cys His Met Leu Ser Arg Asp Thr
        115                 120                 125
Tyr Glu Cys Thr Cys Gln Val Gly Phe Thr Gly Lys Glu Cys Gln Trp
    130                 135                 140
Thr Asp Ala Cys Leu Ser His Pro Cys Ala Asn Gly Ser Thr Cys Thr
145                 150                 155                 160
Thr Val Ala Asn Gln Phe Ser Cys Lys Cys Leu Thr Gly Phe Thr Gly
                165                 170                 175
Gln Lys Cys Glu Thr Asp Val Asn Glu Cys Asp Ile Pro Gly His Cys
            180                 185                 190
Gln His Gly Gly Ile Cys Leu Asn Leu Pro Gly Ser Tyr Gln Cys Gln
        195                 200                 205
Cys Leu Gln Gly Phe Thr Gly Gln Tyr Cys Asp Ser Leu Tyr Val Pro
    210                 215                 220
Cys Ala Pro Ser Pro Cys Val Asn Gly Gly Thr Cys Arg Gln Thr Gly
225                 230                 235                 240
Asp Phe Thr Phe Glu Cys Asn Cys Leu Pro Glu Thr Val Arg Arg Gly
                245                 250                 255
Thr Glu Leu Trp Glu Arg Asp Arg Glu Val Trp Asn Gly Lys Glu His
            260                 265                 270
Asp Glu Asn  *
        275
```

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<210> 1451
<211> 121
<212> PRT
<213> Homo sapiens
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```
<400> 1451
Met Glu Ser Gly Leu Ser Trp Ile Phe Leu Leu Ala Ile Leu Lys Gly
 1               5                  10                  15
Val Gln Cys Glu Val Gln Leu Val Glu Ser Gly Gly Gly Leu Val Gln
            20                  25                  30
Pro Gly Arg Ser Leu Arg Leu Ser Cys Ala Ala Ser Gly Phe Arg Phe
        35                  40                  45
Asp Glu Tyr Gly Met His Trp Val Arg Gln Ala Pro Gly Lys Gly Leu
    50                  55                  60
Glu Trp Val Gly Gly Ile Ser Trp Asn Arg Asp Ser Ile Ala Tyr Ala
65                  70                  75                  80
Asp Ser Val Lys Gly Arg Phe Thr Ile Ser Arg Asp Asn Ala Gln Ser
                85                  90                  95
Tyr Val Tyr Leu Gln Met Asn Ser Leu Arg His Glu Asp Thr Ala Leu
            100                 105                 110
Tyr Tyr Cys Thr Lys Leu Arg Ser Ser
        115                 120 121
```

<210> 1452
<211> 48
<212> PRT
<213> Homo sapiens

<400> 1452
Met Glu Arg Gly Asn Ala Leu Val Val Leu Arg Ser Leu Leu Trp Pro
1 5 10 15
Gly Leu Thr Phe Tyr His Ala Pro Arg Thr Lys Asn Tyr Gly Tyr Val
20 25 30
Tyr Val Gly Thr Gly Glu Lys Asn Met Asp Leu Pro Phe Met Leu *
35 40 45 47

<210> 1453
<211> 123
<212> PRT
<213> Homo sapiens

<400> 1453
Met Ile Thr Val Gln Phe Ser Tyr Thr Ala Val Lys Trp Leu Leu Asn
1 5 10 15
Cys Phe Val Leu Ile Leu Tyr Val Ile Leu Ser Ile Leu Phe Gln Val
20 25 30
Ser Gln Lys Asn Ser Ser Lys Leu Gly Arg Phe Lys Asn Leu Phe Asn
35 40 45
His Lys Glu Cys Ser Lys Leu Leu Phe Asn Arg Asn Gln Ala Gln Thr
50 55 60
Leu Glu Leu Thr Ala Asp Arg Ile Arg Phe Gly Leu Phe Pro Glu Trp
65 70 75 80
Lys His Phe Ser His Thr Thr Ser Leu Cys Thr Ala Lys Met Leu Ala
85 90 95
Tyr Pro Leu Trp Phe Pro Ser Phe Ser Leu Ala Ser Gln Arg Asn Leu
100 105 110
Pro Pro His Pro Leu Tyr Tyr Ile Phe Tyr *
115 120 122

<210> 1454
<211> 327
<212> PRT
<213> Homo sapiens

<400> 1454
Met Arg Glu Trp Trp Val Gln Val Gly Leu Leu Ala Val Pro Leu Leu
1 5 10 15
Ala Ala Tyr Leu His Ile Pro Pro Pro Gln Leu Ser Pro Ala Leu His
20 25 30
Ser Trp Lys Ser Ser Gly Lys Phe Phe Thr Tyr Lys Gly Leu Arg Ile
35 40 45
Phe Tyr Gln Asp Ser Val Gly Val Val Gly Ser Pro Glu Ile Val Val

```
        50                  55                  60
Leu Leu His Gly Phe Pro Thr Ser Ser Tyr Asp Trp Tyr Lys Ile Trp
65                  70                  75                  80
Glu Gly Leu Thr Leu Arg Phe His Arg Val Ile Ala Leu Asp Phe Leu
                85                  90                  95
Gly Phe Gly Phe Ser Asp Lys Pro Arg Pro His His Tyr Ser Ile Phe
            100                 105                 110
Glu Gln Ala Ser Ile Val Glu Ala Leu Leu Arg His Leu Gly Leu Gln
        115                 120                 125
Asn Arg Arg Ile Asn Leu Leu Ser His Asp Tyr Gly Asp Ile Val Ala
    130                 135                 140
Gln Glu Leu Leu Tyr Arg Tyr Lys Gln Asn Arg Ser Gly Arg Leu Thr
145                 150                 155                 160
Ile Lys Ser Leu Cys Leu Ser Asn Gly Gly Ile Phe Pro Glu Thr His
                165                 170                 175
Arg Pro Leu Leu Leu Gln Lys Leu Leu Lys Asp Gly Gly Val Leu Ser
            180                 185                 190
Pro Ile Leu Thr Arg Leu Met Asn Phe Phe Val Phe Ser Arg Gly Leu
        195                 200                 205
Thr Pro Val Phe Gly Pro Tyr Thr Arg Pro Ser Glu Ser Glu Leu Trp
    210                 215                 220
Asp Met Trp Ala Gly Ile Arg Asn Asn Asp Gly Asn Leu Val Ile Asp
225                 230                 235                 240
Ser Leu Leu Gln Tyr Ile Asn Gln Arg Lys Lys Phe Arg Arg Arg Trp
                245                 250                 255
Val Gly Ala Leu Ala Ser Val Thr Ile Pro Ile His Phe Ile Tyr Gly
            260                 265                 270
Pro Leu Asp Pro Val Asn Pro Tyr Pro Glu Phe Leu Glu Leu Tyr Arg
        275                 280                 285
Lys Thr Leu Pro Arg Ser Thr Val Ser Ile Leu Asp Asp His Ile Ser
    290                 295                 300
His Tyr Pro Gln Leu Glu Asp Pro Met Gly Phe Leu Asn Ala Tyr Met
305                 310                 315                 320
Gly Phe Ile Asn Ser Phe  *
                325 326


     <210> 1455
     <211> 57
     <212> PRT
     <213> Homo sapiens


     <400> 1455
Met Ile Leu Leu Lys Val Cys Ser Ala Ala Ser Leu Leu Gly Glu Gly
1               5                   10                  15
Phe Met Asn Gln Val Thr Ser Thr Asn Lys Ala Ser Leu Ser Leu Leu
            20                  25                  30
Ser Leu Thr Met Lys Val Ala Val Asn Lys Gly Lys Lys Glu Arg Glu
        35                  40                  45
Leu Phe Ile Pro Phe Gln Phe Gln  *
    50                  55  56


     <210> 1456
     <211> 48
     <212> PRT
```

<213> Homo sapiens

<400> 1456
```
Met His Cys Ile Phe Ser Cys Leu Leu Trp Cys Ile Gln Leu Pro Ser
 1               5                  10                  15
Met Leu Ser Val Leu Lys Thr Gln Pro Ser Lys Asn His Pro Leu Trp
            20                  25                  30
Pro Cys Lys Tyr Ala Tyr Asn Ile Phe Phe Phe Leu Cys Ile Ile  *
        35                  40                  45      47
```

<210> 1457
<211> 459
<212> PRT
<213> Homo sapiens

<400> 1457
```
Met Ser Asp Leu Leu Ser Val Phe Leu His Leu Leu Leu Leu Phe Lys
 1               5                  10                  15
Leu Val Ala Pro Val Thr Phe Arg His His Arg Tyr Asp Asp Leu Val
            20                  25                  30
Arg Thr Leu Tyr Lys Val Gln Asn Glu Cys Pro Gly Ile Thr Arg Val
        35                  40                  45
Tyr Ser Ile Gly Arg Ser Val Glu Gly Arg His Leu Tyr Val Leu Glu
    50                  55                  60
Phe Ser Asp His Pro Gly Ile His Glu Pro Leu Glu Pro Glu Val Lys
65                  70                  75                  80
Tyr Val Gly Asn Met His Gly Asn Glu Ala Leu Gly Arg Glu Leu Met
                85                  90                  95
Leu Gln Leu Ser Glu Phe Leu Cys Glu Glu Phe Arg Asn Arg Asn Gln
            100                 105                 110
Arg Ile Val Gln Leu Ile Gln Asp Thr Arg Ile His Ile Leu Pro Ser
        115                 120                 125
Met Asn Pro Asp Gly Tyr Glu Val Ala Ala Ala Gln Gly Pro Asn Lys
    130                 135                 140
Pro Gly Tyr Leu Val Gly Arg Asn Asn Ala Asn Gly Val Asp Leu Asn
145                 150                 155                 160
Arg Asn Phe Pro Asp Leu Asn Thr Tyr Ile Tyr Tyr Asn Glu Lys Tyr
                165                 170                 175
Gly Gly Pro Asn His His Leu Pro Leu Pro Asp Asn Trp Lys Ser Gln
            180                 185                 190
Val Glu Pro Glu Thr Arg Ala Val Ile Arg Trp Met His Ser Phe Asn
        195                 200                 205
Phe Val Leu Ser Ala Asn Leu His Gly Gly Ala Val Val Ala Asn Tyr
    210                 215                 220
Pro Tyr Asp Lys Ser Phe Glu His Arg Val Arg Gly Val Arg Arg Thr
225                 230                 235                 240
Ala Ser Thr Pro Thr Pro Asp Asp Lys Leu Phe Gln Lys Leu Ala Lys
                245                 250                 255
Val Tyr Ser Tyr Ala His Gly Trp Met Phe Gln Gly Trp Asn Cys Gly
            260                 265                 270
Asp Tyr Phe Pro Asp Gly Ile Thr Asn Gly Ala Ser Trp Tyr Ser Leu
        275                 280                 285
Ser Lys Gly Met Gln Asp Phe Asn Tyr Leu His Thr Asn Cys Phe Glu
    290                 295                 300
Ile Thr Leu Glu Leu Ser Cys Asp Lys Phe Pro Pro Glu Glu Glu Leu
```

```
305                 310                 315                 320
Gln Arg Glu Trp Leu Gly Asn Arg Glu Ala Leu Ile Gln Phe Leu Glu
                325                 330                 335
Gln Val His Gln Gly Ile Lys Gly Met Val Leu Asp Glu Asn Tyr Asn
            340                 345                 350
Asn Leu Ala Asn Ala Val Ile Ser Val Ser Gly Ile Asn His Asp Val
        355                 360                 365
Thr Ser Gly Asp His Gly Asp Tyr Phe Arg Leu Leu Leu Pro Gly Ile
    370                 375                 380
Tyr Thr Val Ser Ala Thr Ala Pro Gly Tyr Asp Pro Glu Thr Val Thr
385                 390                 395                 400
Val Thr Val Gly Pro Ala Glu Pro Thr Leu Val Asn Phe His Leu Lys
                405                 410                 415
Arg Ser Ile Pro Gln Val Ser Pro Val Arg Arg Ala Pro Ser Arg Arg
            420                 425                 430
His Gly Val Arg Ala Lys Val Gln Pro Gln Pro Arg Lys Lys Glu Met
        435                 440                 445
Glu Met Arg Gln Leu Gln Arg Gly Pro Ala  *
    450                 455         458
```

```
<210> 1458
<211> 463
<212> PRT
<213> Homo sapiens
```

```
<400> 1458
Met Ala Arg Val Leu Gly Ala Pro Val Ala Leu Gly Leu Trp Ser Leu
 1               5                  10                  15
Cys Trp Ser Leu Ala Ile Ala Thr Pro Leu Pro Pro Thr Ser Ala His
            20                  25                  30
Gly Asn Val Ala Glu Gly Glu Thr Lys Pro Asp Pro Asp Val Thr Glu
        35                  40                  45
Arg Cys Ser Asp Gly Trp Ser Phe Asp Ala Thr Thr Leu Asp Asp Asn
    50                  55                  60
Gly Thr Met Leu Phe Phe Lys Gly Glu Phe Val Trp Lys Ser His Lys
65                  70                  75                  80
Trp Asp Arg Glu Leu Ile Ser Glu Arg Trp Lys Asn Phe Pro Ser Pro
                85                  90                  95
Val Asp Ala Ala Phe Arg Gln Gly His Asn Ser Val Phe Leu Ile Lys
            100                 105                 110
Gly Asp Lys Val Trp Val Tyr Pro Pro Glu Lys Lys Glu Lys Gly Tyr
        115                 120                 125
Pro Lys Leu Leu Gln Asp Glu Phe Pro Gly Ile Pro Ser Pro Leu Asp
    130                 135                 140
Ala Ala Val Glu Cys His Arg Gly Glu Cys Gln Ala Glu Gly Val Leu
145                 150                 155                 160
Phe Phe Gln Gly Asp Arg Glu Trp Phe Trp Asp Leu Ala Thr Gly Thr
                165                 170                 175
Met Lys Glu Arg Ser Trp Pro Ala Val Gly Asn Cys Ser Ser Ala Leu
            180                 185                 190
Arg Trp Leu Gly Arg Tyr Tyr Cys Phe Gln Gly Asn Gln Phe Leu Arg
        195                 200                 205
Phe Asp Pro Val Arg Gly Glu Val Pro Pro Arg Tyr Pro Arg Asp Val
    210                 215                 220
Arg Asp Tyr Phe Met Pro Cys Pro Gly Arg Gly His Gly His Arg Asn
225                 230                 235                 240
```

Gly Thr Gly His Gly Asn Ser Thr His His Gly Pro Glu Tyr Met Arg
245 250 255
Cys Ser Pro His Leu Val Leu Ser Ala Leu Thr Ser Asp Asn His Gly
260 265 270
Ala Thr Tyr Ala Phe Ser Gly Thr His Tyr Trp Arg Leu Asp Thr Ser
275 280 285
Arg Asp Gly Trp His Ser Trp Pro Ile Ala His Gln Trp Pro Gln Gly
290 295 300
Pro Ser Ala Val Asp Ala Ala Phe Ser Trp Glu Glu Lys Leu Tyr Leu
305 310 315 320
Val Gln Gly Thr Gln Val Tyr Val Phe Leu Thr Lys Gly Gly Tyr Thr
325 330 335
Leu Val Ser Gly Tyr Pro Lys Arg Leu Glu Lys Glu Val Gly Thr Pro
340 345 350
His Gly Ile Ile Leu Asp Ser Val Asp Ala Ala Phe Ile Cys Pro Gly
355 360 365
Ser Ser Arg Leu His Ile Met Ala Gly Arg Arg Leu Trp Trp Leu Asp
370 375 380
Leu Lys Ser Gly Ala Gln Ala Thr Trp Thr Glu Leu Pro Trp Pro His
385 390 395 400
Glu Lys Val Asp Gly Ala Leu Cys Met Glu Lys Ser Leu Gly Pro Asn
405 410 415
Ser Cys Ser Ala Asn Gly Pro Gly Leu Tyr Leu Ile His Gly Pro Asn
420 425 430
Leu Tyr Cys Tyr Ser Asp Val Glu Lys Leu Asn Ala Ala Lys Ala Leu
435 440 445
Pro Gln Pro Gln Asn Val Thr Ser Leu Leu Gly Cys Thr His *
450 455 460 462

<210> 1459
<211> 187
<212> PRT
<213> Homo sapiens

<400> 1459
Met Gln Pro Ile Val Ala Lys Ala Leu Val Val Leu Leu Glu Val His
1 5 10 15
Pro Leu Gln Asp Gln Ala Glu Ser Gly Arg Leu Gly His Val His Leu
20 25 30
Leu Cys Ala Pro Ala Ala Leu Gln His Ala Leu Arg Gly Ile Thr Leu
35 40 45
His Asn Gly His His Gln Ala Asp His Leu Pro Asp Leu Met His His
50 55 60
Glu Ala Leu Ala Leu His Pro Asp His Arg Lys Leu Gln Ala Leu Pro
65 70 75 80
His Lys Gly Phe Leu Ala Val His Leu Gln Asp Val Ala Ala Gly Thr
85 90 95
Gly Ile Leu Arg Pro Leu Leu Arg Gly Glu Ile Val Glu Val Val Arg
100 105 110
Ala Leu Val Ala Gly Gln Glu Pro Val Asp Leu Leu Gln Arg Leu Gly
115 120 125
Ala Gln Ala Val Gly Leu Ile Leu Asn Val Pro Val Leu Val Arg Lys
130 135 140
Gly Lys Arg Gly Gln Gln Val Ala Ile Gly Pro Gly Ile Thr Ser Val
145 150 155 160
Leu Gly Val Lys Pro Ala Arg Asp Pro Leu Gln Ser Gln Asn Pro Asn

```
                165                 170                 175
Val Arg Gly Lys Val Ala Val Asp Leu Phe  *
            180                 185 186
```

<210> 1460
<211> 223
<212> PRT
<213> Homo sapiens

<400> 1460

```
Met Lys Phe Ala Leu Phe Thr Ser Gly Val Ala Leu Thr Leu Ser Phe
  1               5                  10                  15
Val Phe Met Tyr Ala Lys Cys Glu Asn Glu Pro Phe Ala Gly Val Ser
            20                  25                  30
Glu Ser Tyr Asn Gly Thr Gly Glu Leu Gly Asn Leu Ile Ala Pro Cys
        35                  40                  45
Asn Ala Asn Cys Asn Cys Ser Arg Ser Tyr Tyr Tyr Pro Val Cys Gly
    50                  55                  60
Asp Gly Val Gln Tyr Phe Ser Pro Cys Phe Ala Gly Cys Ser Asn Pro
 65                  70                  75                  80
Val Ala His Arg Lys Pro Lys Val Tyr Tyr Asn Cys Ser Cys Ile Glu
                 85                  90                  95
Arg Lys Thr Glu Ile Thr Ser Thr Ala Glu Thr Phe Gly Phe Glu Ala
            100                 105                 110
Asn Ala Gly Lys Cys Glu Thr His Cys Ala Lys Leu Ala Ile Phe Leu
        115                 120                 125
Cys Ile Val Phe Ile Gly Asn Ile Phe Thr Phe Met Ala Arg Ser Pro
    130                 135                 140
Ile Thr Gly Ala Ile Pro Arg Gly Gly Asn His Arg Gln Arg Pro Pro
145                 150                 155                 160
Thr Leu Gly Ile Gln Phe Met Ala Leu Arg Thr Leu Trp Thr Thr Pro
                165                 170                 175
Trp Pro Ser Lys Thr Gly Cys Pro Ile His Gln Pro Gly Ser Leu Trp
            180                 185                 190
Glu Lys Leu Gly Trp Arg Pro Leu Lys Thr Leu Arg Arg Pro Lys Pro
        195                 200                 205
Ser Trp Asn Ala Leu Leu Ala Leu Ala His Pro Arg Ser Phe Gln
    210                 215                 220         223
```

<210> 1461
<211> 210
<212> PRT
<213> Homo sapiens

<400> 1461

```
Met Tyr Phe Phe Leu Leu Leu Leu Phe Phe Asn Val Gln Arg Leu Ala
  1               5                  10                  15
Phe Pro Phe Gly Ile Pro Asn Asp Pro Met Leu Trp Ser Glu Gly Gln
            20                  25                  30
Ser His Leu Cys Trp Arg Ser Pro Leu Ile Pro Ser Ala Gln Phe Arg
        35                  40                  45
Gly Ser Arg Ala Asp Ile Arg Gly Ser Met Leu His Ser Ser Ser Gly
    50                  55                  60
```

Arg Val Val Pro Leu Asn Pro Ala Thr Lys Leu Ser Pro Leu Glu Ser
65 70 75 80
Gln Met Ala Leu His Thr Lys Ala Val Glu Ala Gly Met Val Phe Gly
85 90 95
His Arg Ala Glu His Lys Asp Pro Arg Ser Val Trp Glu Ser Tyr Trp
100 105 110
Leu Leu Gly Ser Pro Trp Ala Glu Val Thr Arg Leu His Pro Arg Arg
115 120 125
Ala Gln Leu Gly Ser Leu Pro Pro Pro Asp Pro Arg Thr Thr His Arg
130 135 140
Arg Gly Ala Val Ser Ile Phe Leu Lys Gly Pro Phe Gly Asp Leu Val
145 150 155 160
Leu Ser Val Glu Arg Thr Asp Val Ala Leu Ser Ser Gln His Ile Pro
165 170 175
Gly Ser Gly Arg Pro Gln Leu Lys Gln Cys Gln Gly Pro Gln Gly Ser
180 185 190
His Leu Asp Arg Pro Thr Ala Cys Asn Ser Ala Leu Leu Arg Arg Gln
195 200 205
His *
209

<210> 1462
<211> 56
<212> PRT
<213> Homo sapiens

<400> 1462
Met Ala Val Arg Val Leu Trp Gly Gly Leu Ser Leu Leu Arg Val Leu
1 5 10 15
Trp Cys Leu Leu Pro Gln Thr Gly Tyr Val His Pro Asp Glu Phe Phe
20 25 30
Gln Ser Pro Glu Val Met Ala Gly Lys Thr Pro His Val Trp Leu Arg
35 40 45
Gln Ala Ala Ala Glu Ser Ala *
50 55

<210> 1463
<211> 66
<212> PRT
<213> Homo sapiens

<400> 1463
Met Glu Asn Cys Val Gly Glu Arg Asn His Pro Leu Phe Val Val Tyr
1 5 10 15
Leu Ala Leu Gln Leu Val Val Leu Leu Trp Gly Leu Tyr Leu Ala Cys
20 25 30
Pro Gly Val Cys Gly Cys Gly Pro Ala Gly Ser Cys Ser Pro Pro Ser
35 40 45
Cys Cys Trp Pro Ser Ser Arg Gly Gly Gln Pro Gly Ser Arg Leu Ala
50 55 60
Pro Leu
65 66

<210> 1464
<211> 200
<212> PRT
<213> Homo sapiens

<400> 1464
```
Met Val Trp Arg Arg Leu Leu Arg Lys Arg Trp Val Leu Ala Leu Val
 1               5                  10                  15
Phe Gly Leu Ser Leu Val Tyr Phe Leu Ser Ser Thr Phe Lys Gln Glu
            20                  25                  30
Glu Arg Ala Val Arg Asp Arg Asn Leu Leu Gln Val His Asp His Asn
        35                  40                  45
Gln Pro Ile Pro Trp Lys Val Gln Phe Asn Leu Gly Asn Ser Ser Arg
    50                  55                  60
Pro Ser Asn Gln Cys Arg Asn Ser Ile Gln Gly Lys His Leu Ile Thr
65                  70                  75                  80
Asp Glu Leu Gly Tyr Val Cys Glu Arg Lys Asp Leu Leu Val Asn Gly
                85                  90                  95
Cys Cys Asn Val Asn Val Pro Ser Thr Lys Gln Tyr Cys Cys Asp Gly
            100                 105                 110
Cys Trp Pro Asn Gly Cys Cys Ser Ala Tyr Glu Tyr Cys Val Ser Cys
        115                 120                 125
Cys Leu Gln Pro Asn Lys Gln Leu Leu Leu Glu Arg Phe Leu Asn Arg
    130                 135                 140
Ala Ala Val Ala Phe Gln Asn Leu Phe Met Ala Val Glu Asp His Phe
145                 150                 155                 160
Glu Leu Cys Leu Ala Lys Cys Arg Thr Ser Ser Gln Ser Val Gln His
                165                 170                 175
Glu Asn Thr Tyr Arg Asp Pro Ile Ala Lys Tyr Cys Tyr Gly Glu Ser
            180                 185                 190
Pro Pro Glu Leu Phe Pro Ala  *
        195             199
```

<210> 1465
<211> 46
<212> PRT
<213> Homo sapiens

<400> 1465
```
Met Gln Leu Ile Arg Arg Ser His Asn Arg His Trp Phe Arg Ser Ala
 1               5                  10                  15
Ile Thr Phe Leu Met Cys Lys Gly Ile Thr Leu Leu Trp Leu Trp Lys
            20                  25                  30
Leu Leu Thr Gly Asn Asp Cys Ile Glu Tyr Ile Arg Lys  *
        35                  40                  45
```

<210> 1466
<211> 56
<212> PRT
<213> Homo sapiens

<400> 1466
Met Arg Leu Leu Phe Ser Ser Gln Val Asn Ser Lys Arg Leu Thr Ala
1 5 10 15
Ser Arg Ala Phe Leu Val Leu Val Pro Ala His Leu Ser Tyr Leu Leu
20 25 30
Ala Leu Pro Ser Ile Pro Ala Thr Arg Gly Phe Trp Phe Lys Asp Thr
35 40 45
Val Phe Leu Ser Cys Ser Ala *
50 55

<210> 1467
<211> 366
<212> PRT
<213> Homo sapiens

<400> 1467
Met Arg Gly Gln Val Val Thr Leu Ile Leu Leu Leu Leu Leu Lys Val
1 5 10 15
Tyr Gln Gly Lys Gly Cys Gln Gly Ser Ala Asp His Val Val Ser Ile
20 25 30
Ser Gly Val Pro Leu Gln Leu Gln Pro Asn Ser Ile Gln Thr Lys Val
35 40 45
Asp Ser Ile Ala Trp Lys Lys Leu Leu Pro Ser Gln Asn Gly Phe His
50 55 60
His Ile Leu Lys Trp Glu Asn Gly Ser Leu Pro Ser Asn Thr Ser Asn
65 70 75 80
Asp Arg Phe Ser Phe Ile Val Lys Asn Leu Ser Leu Leu Ile Lys Ala
85 90 95
Ala Gln Gln Gln Asp Ser Gly Leu Tyr Cys Leu Glu Val Thr Ser Ile
100 105 110
Ser Gly Lys Val Gln Thr Ala Thr Phe Gln Val Phe Val Phe Asp Lys
115 120 125
Val Glu Lys Pro Arg Leu Gln Gly Gln Gly Lys Ile Leu Asp Arg Gly
130 135 140
Arg Cys Gln Val Ala Leu Ser Cys Leu Val Ser Arg Asp Gly Asn Val
145 150 155 160
Ser Tyr Ala Trp Tyr Arg Gly Ser Lys Leu Ile Gln Thr Ala Gly Asn
165 170 175
Leu Thr Tyr Leu Asp Glu Glu Val Asp Ile Asn Gly Thr His Thr Tyr
180 185 190
Thr Cys Asn Val Ser Asn Pro Val Ser Trp Glu Ser His Thr Leu Asn
195 200 205
Leu Thr Gln Asp Cys Gln Asn Ala His Gln Glu Phe Arg Phe Trp Pro
210 215 220
Phe Leu Val Ile Ile Val Ile Leu Ser Ala Leu Phe Leu Gly Thr Leu
225 230 235 240
Ala Cys Phe Cys Val Trp Arg Arg Lys Arg Lys Glu Lys Gln Ser Glu
245 250 255
Thr Ser Pro Lys Glu Phe Leu Thr Ile Tyr Glu Asp Val Lys Asp Leu
260 265 270
Lys Thr Arg Arg Asn His Glu Gln Glu Gln Thr Phe Pro Gly Gly Gly
275 280 285
Ser Thr Ile Tyr Ser Met Ile Gln Ser Gln Ser Ser Ala Pro Thr Ser
290 295 300
Gln Glu Pro Ala Tyr Thr Leu Tyr Ser Leu Ile Gln Pro Ser Arg Lys

```
305                 310                 315                 320
Ser Gly Ser Arg Lys Arg Asn His Ser Pro Ser Phe Asn Ser Thr Ile
                325                 330                 335
Tyr Glu Val Ile Gly Lys Ser Gln Pro Lys Ala Gln Asn Pro Ala Arg
            340                 345                 350
Leu Ser Arg Lys Glu Leu Glu Asn Phe Asp Val Tyr Ser  *
        355                 360                 365
```

```
     <210> 1468
     <211> 57
     <212> PRT
     <213> Homo sapiens


     <400> 1468
Met Thr Asp Phe Phe Leu Cys Ile His Ser Phe Tyr Leu Cys Val Leu
  1               5                  10                  15
Leu Gln Ala Ser Leu Asp Met Leu Ser Val Lys Ser Phe Ser Phe Lys
             20                  25                  30
Val Leu Cys Leu Met Lys Ala Lys Glu Lys Pro Asn Thr Thr Ser Cys
         35                  40                  45
His Leu Val Ile Asp Ser Asn Ser Thr
     50                  55      57
```

```
     <210> 1469
     <211> 110
     <212> PRT
     <213> Homo sapiens


     <400> 1469
Met Leu Glu Ile Leu Leu Lys Leu Val Arg Leu Leu Thr Thr Gln Pro
  1               5                  10                  15
Tyr Leu Thr Leu Phe Gln Ala Val Arg Asn Leu Ala Leu Asn Leu Ser
             20                  25                  30
Thr Ser Ser Gly Ser Leu Gly Pro Ala Pro Gly Glu Pro Arg Ala Gly
         35                  40                  45
Pro Leu Ala Pro Glu Gly Pro Arg Pro Leu Gly Ser Gly Pro Leu Gly
     50                  55                  60
Pro Arg Gly Leu Arg Ala Ser Gly Arg Arg Arg Ala Ser Ser Gly Leu
 65                  70                  75                  80
Leu Leu Arg Tyr Cys Ala Ala Ala Gly Asp Thr Glu Phe Met Asp Ala
                 85                  90                  95
Pro Gly Gly Arg Thr Glu Gly Pro Gly Gly Gly Leu Arg Pro
            100                 105                 110
```

```
     <210> 1470
     <211> 59
     <212> PRT
     <213> Homo sapiens


     <400> 1470
```

```
Met Met Cys Arg Cys Met Cys Ala Cys Val Cys Ala Pro Val Cys Val
 1               5                  10                  15
His Met His Gly Leu Ala Pro Ala Pro Ala Ile Trp Ile Glu Gln Phe
            20                  25                  30
Trp Val Glu Asn Phe Phe Ser Pro Phe Leu Lys Val Ser Phe Tyr Ser
        35                  40                  45
Leu Pro Val Cys Ile Glu Lys Ser Ser Ile  *
    50                  55          58
```

<210> 1471
<211> 123
<212> PRT
<213> Homo sapiens

<400> 1471

```
Met Met His Phe Leu Thr Gly Gly Trp Lys Val Leu Phe Ala Cys Val
 1               5                  10                  15
Pro Pro Thr Glu Tyr Cys His Gly Trp Ala Cys Phe Gly Val Ser Ile
            20                  25                  30
Leu Val Ile Gly Leu Leu Thr Ala Leu Ile Gly Asp Leu Ala Ser His
        35                  40                  45
Phe Gly Cys Thr Val Gly Leu Lys Asp Ser Val Asn Ala Val Val Phe
    50                  55                  60
Val Ala Leu Gly Thr Ser Ile Pro Gly Asn Thr Leu Gly Asp Phe Gly
65                  70                  75                  80
Gly Val Gly Ser Gln Met Ser Gln Ala Gly Ala Thr Gln Asp Pro Ala
                85                  90                  95
Glu Met Arg His Val Arg Gln Gln Gly Gly Gly Ala Ala Gly Pro Val
            100                 105                 110
Arg Arg Arg Val His Arg Glu Arg Asp Pro Leu
        115                 120         123
```

<210> 1472
<211> 316
<212> PRT
<213> Homo sapiens

<400> 1472

```
Met Val Ser Ala Ser Gly Thr Ser Phe Phe Lys Gly Met Leu Leu Gly
 1               5                  10                  15
Ser Ile Ser Trp Val Leu Ile Thr Met Phe Gly Gln Ile His Ile Arg
            20                  25                  30
His Arg Gly Gln Thr Gln Asp His Glu His His His Leu Arg Pro Pro
        35                  40                  45
Asn Arg Asn Asp Phe Leu Asn Thr Ser Lys Val Ile Leu Leu Glu Leu
    50                  55                  60
Ser Lys Ser Ile Arg Val Phe Cys Ile Ile Phe Gly Glu Ser Glu Asp
65                  70                  75                  80
Glu Ser Tyr Trp Ala Val Leu Lys Glu Thr Trp Thr Lys His Cys Asp
                85                  90                  95
Lys Ala Glu Leu Tyr Asp Thr Lys Asn Asp Asn Leu Phe Asn Ile Glu
            100                 105                 110
Ser Asn Asp Arg Trp Val Gln Met Arg Thr Ala Tyr Lys Tyr Val Phe
```

```
        115                 120                 125
Glu Lys Asn Gly Asp Asn Tyr Asn Trp Phe Phe Leu Ala Leu Pro Thr
    130                 135                 140
Thr Phe Ala Val Ile Glu Asn Leu Lys Tyr Leu Leu Phe Thr Arg Asp
145                 150                 155                 160
Ala Ser Gln Pro Phe Tyr Leu Gly His Thr Val Ile Phe Gly Asp Leu
                165                 170                 175
Glu Tyr Val Thr Val Glu Gly Gly Ile Val Leu Ser Arg Glu Leu Met
            180                 185                 190
Lys Arg Leu Asn Arg Leu Leu Asp Asn Ser Glu Thr Cys Ala Asp Gln
        195                 200                 205
Ser Val Ile Trp Lys Leu Ser Glu Asp Lys Gln Leu Ala Ile Cys Leu
    210                 215                 220
Lys Tyr Ala Gly Val His Ala Glu Asn Ala Glu Asp Tyr Glu Gly Arg
225                 230                 235                 240
Asp Val Phe Asn Thr Lys Pro Ile Ala Gln Leu Ile Glu Glu Ala Leu
                245                 250                 255
Ser Asn Asn Pro Gln Gln Val Val Glu Gly Cys Cys Ser Asp Met Ala
            260                 265                 270
Ile Thr Phe Asn Gly Leu Thr Pro Gln Lys Met Glu Val Met Met Tyr
        275                 280                 285
Gly Leu Tyr Arg Leu Arg Ala Phe Gly His Tyr Phe Asn Asp Thr Leu
    290                 295                 300
Val Phe Leu Pro Pro Val Gly Ser Glu Asn Asp  *
305                 310                 315


     <210> 1473
     <211> 65
     <212> PRT
     <213> Homo sapiens


     <400> 1473
Met Gln Cys Pro Pro Pro Phe Leu Gly Gln Trp Leu Leu Cys Pro Ala
 1               5                  10                  15
Ala Arg Gln Trp Gly Pro Gly Ala Gly Ser Pro Gly Pro Val Leu Val
            20                  25                  30
Pro Ala Gly Arg Arg Arg Pro Pro Pro Arg Ser Gly Pro Gln Arg Asp
        35                  40                  45
Ser Pro Ala Pro Val Arg Gly Pro Gln Phe His Ser Val Val Gly Pro
    50                  55                  60                  64
 *


     <210> 1474
     <211> 55
     <212> PRT
     <213> Homo sapiens


     <400> 1474
Met Ile Phe Met Arg Val Leu Met Leu Leu Cys Cys Met Asp Ser Leu
 1               5                  10                  15
Gly Ser Leu Asp Thr Phe Gln Trp Leu Ser Arg Val Leu Cys Pro Thr
            20                  25                  30
```

```
Glu Asn Leu Ile Phe Glu Leu Asn Gly Tyr Glu Leu Asn Ser Thr Trp
        35                  40                  45
Phe Gly Trp Leu Asn Thr  *
    50              54
```

<210> 1475
<211> 128
<212> PRT
<213> Homo sapiens

<221> misc_feature
<222> (1)...(128)
<223> Xaa = any amino acid or nothing

<400> 1475

```
Met Lys Phe Gln Leu Phe Leu Ser Tyr Val Phe Ile Thr Gln Val Phe
  1               5                  10                  15
Ser Arg Pro Phe Gln Ser Asn Leu Gly Ser Leu Thr Pro Ala Ser Ser
            20                  25                  30
Gln Ile Pro Leu Gln Leu Pro Lys Ala Leu Cys Val Arg Cys Leu Asn
        35                  40                  45
Thr Val Xaa Xaa Xaa Xaa Xaa Thr Gly Phe Gly Lys Phe Gln Ile Thr
    50                  55                  60
Ile Gln Ser Pro Gly Gly Pro Leu Val Leu Ala Arg Pro Trp Ala Ser
 65                  70                  75                  80
Lys Phe Pro Ser Pro Lys Phe Xaa Xaa Xaa Xaa Xaa Xaa Pro Lys Met
                 85                  90                  95
Gly Gly Lys Thr Phe Ala Tyr Gly Arg Ile Asn Pro Thr Arg Pro Ala
            100                 105                 110
Lys Asn Xaa Xaa Xaa Xaa Xaa Xaa Ser Leu Ala Ser Leu Asn Pro Thr
        115                 120                 125         128
```

<210> 1476
<211> 210
<212> PRT
<213> Homo sapiens

<400> 1476

```
Met Tyr Phe Phe Leu Leu Leu Leu Phe Phe Asn Val Gln Arg Leu Ala
  1               5                  10                  15
Phe Pro Phe Gly Ile Pro Asn Asp Pro Met Leu Trp Ser Glu Gly Gln
            20                  25                  30
Ser His Leu Cys Trp Arg Ser Pro Leu Ile Pro Ser Ala Gln Phe Arg
        35                  40                  45
Gly Ser Arg Ala Asp Ile Arg Gly Ser Met Leu His Ser Ser Ser Gly
    50                  55                  60
Arg Val Val Pro Leu Asn Pro Ala Thr Lys Leu Ser Pro Leu Glu Ser
 65                  70                  75                  80
Gln Met Ala Leu His Thr Lys Ala Val Glu Ala Gly Met Val Phe Gly
                 85                  90                  95
His Arg Ala Glu His Lys Asp Pro Arg Ser Val Trp Glu Ser Tyr Trp
```

```
            100                 105                 110
Leu Leu Gly Ser Pro Trp Ala Glu Val Thr Arg Leu His Pro Arg Arg
        115                 120                 125
Ala Gln Leu Gly Ser Leu Pro Pro Pro Asp Pro Arg Thr Thr His Arg
    130                 135                 140
Arg Gly Ala Val Ser Ile Phe Leu Lys Gly Pro Phe Gly Asp Leu Val
145                 150                 155                 160
Leu Ser Val Glu Arg Thr Asp Val Ala Leu Ser Ser Gln His Ile Pro
                165                 170                 175
Gly Ser Gly Arg Pro Gln Leu Lys Gln Cys Gln Gly Pro Gln Gly Ser
            180                 185                 190
His Leu Asp Arg Pro Thr Ala Cys Asn Ser Ala Leu Leu Arg Arg Gln
        195                 200                 205
His  *
209
```

```
     <210> 1477
     <211> 57
     <212> PRT
     <213> Homo sapiens


     <400> 1477
Met His Thr Cys Gln Ile Tyr Ile Tyr Ser Thr Asn Val Thr Phe Leu
  1               5                  10                  15
Phe Phe Val Leu Asp Val Arg Ala Cys Ser Tyr Val Arg Tyr Leu His
            20                  25                  30
Lys Leu Leu His Tyr Phe Phe Leu Cys Asn Thr Phe Leu Phe Val Tyr
        35                  40                  45
Val Val Gln Ile Tyr Phe Phe Pro  *
     50                  55  56
```

```
     <210> 1478
     <211> 97
     <212> PRT
     <213> Homo sapiens


     <400> 1478
Met Arg Ile Trp Ser Arg Ala Val Gly Asp Gly Pro Ala Ala Val Cys
  1               5                  10                  15
Cys Pro Leu Arg Ser Trp Cys Leu Leu Leu Trp Ala Leu Asp Ser Leu
            20                  25                  30
Asp Pro Ala Ala Val Thr Thr His Ala Ser Ala Met Leu Ser Gly Val
        35                  40                  45
Phe Thr Pro Pro Phe Val Ser Ala Leu Pro Val Gln Trp Met Gln Met
     50                  55                  60
Pro Val Leu Ser Phe Leu Ser Leu Thr Gly Ser Ser Val Tyr Val His
 65                  70                  75                  80
Met Ala Leu Leu Ser Gly His Gln Gly Ser Asp Thr Cys Ser Gly Leu
                 85                  90                  95  96
 *
```

<210> 1479
<211> 113
<212> PRT
<213> Homo sapiens

<400> 1479
```
Met Leu Ser Ile Ser Tyr Phe Ser Asn Ser Leu Met Leu Arg Leu Val
 1               5                  10                  15
Pro Leu Ala Ala Tyr Val Leu Ser Tyr Leu Ile Cys Ser Val Leu Leu
            20                  25                  30
His Ile Asn Gln Thr Thr Val Thr Thr Tyr Arg Gly Arg Lys Gln Arg
        35                  40                  45
Lys Lys Ile Gln Phe Ala Thr Gly Asn His Gln Ser Ala Gln Ser Tyr
    50                  55                  60
Ser Glu Leu Leu Ser Leu Ser Leu Ser Phe Ser Ser Leu Leu Ser Pro
65                  70                  75                  80
Val Phe Ser Leu Pro Ser Trp Ser Leu Pro Ser Leu Pro Pro Phe Phe
                85                  90                  95
Ser His Ser Pro His Gln Lys Gly Ile Met Met Val Pro Arg Ser Val
            100                 105                 110     112
 *
```

<210> 1480
<211> 91
<212> PRT
<213> Homo sapiens

<400> 1480
```
Met Arg Leu Ser Val Cys Leu Leu Leu Leu Thr Leu Ala Leu Cys Cys
 1               5                  10                  15
Tyr Arg Ala Asn Ala Val Val Cys Gln Ala Leu Gly Ser Glu Ile Thr
            20                  25                  30
Gly Phe Leu Leu Ala Gly Lys Pro Val Phe Lys Phe Gln Leu Ala Lys
        35                  40                  45
Phe Lys Ala Pro Leu Glu Ala Val Ala Ala Lys Met Glu Val Lys Lys
    50                  55                  60
Cys Val Asp Thr Met Ala Tyr Glu Lys Arg Val Leu Ile Thr Lys Thr
65                  70                  75                  80
Leu Gly Lys Ile Ala Glu Lys Cys Asp Arg  *
                85                  90
```

<210> 1481
<211> 54
<212> PRT
<213> Homo sapiens

<400> 1481
```
Met Pro Gly Ser Ile Leu Ser Asn Leu His Val Leu Leu Lys Tyr Leu
 1               5                  10                  15
Phe Thr Phe Ala Glu Val Phe Leu Val Pro Gly Pro Phe Asn Val Leu
```

```
            20                  25                  30
Phe Leu Ser Leu Arg Leu Glu Thr Leu Thr Phe Phe Val Leu Trp Leu
        35                  40                  45
Val Pro Tyr Leu Ile  *
    50          53
```

<210> 1482
<211> 56
<212> PRT
<213> Homo sapiens

<400> 1482

```
Met Glu Arg Trp Leu Gly Leu Ile Gln Thr Leu Trp Leu Pro Ala His
 1               5                  10                  15
Ser Gly Pro Leu Gly Arg Ala Trp Val Val Pro Arg Ala Thr Ser Gly
            20                  25                  30
His Tyr Trp Gly Gly Lys Gly Thr Asn Glu Gly Gly Gln Asp Lys Gly
        35                  40                  45
His Phe Pro Leu Pro Pro Arg  *
    50                  55
```

<210> 1483
<211> 202
<212> PRT
<213> Homo sapiens

<400> 1483

```
Met Leu Leu Leu Leu Gly Leu Cys Leu Gly Leu Ser Leu Cys Val Gly
 1               5                  10                  15
Ser Gln Glu Glu Ala Gln Ser Trp Gly His Ser Ser Glu Gln Asp Gly
            20                  25                  30
Leu Arg Val Pro Arg Gln Val Arg Leu Leu Gln Arg Leu Lys Thr Lys
        35                  40                  45
Pro Leu Met Thr Glu Phe Ser Val Lys Ser Thr Ile Ile Ser Arg Tyr
    50                  55                  60
Ala Phe Thr Thr Val Ser Cys Arg Met Leu Asn Arg Ala Ser Glu Asp
65                  70                  75                  80
Gln Asp Ile Glu Phe Gln Met Gln Ile Pro Ala Ala Ala Phe Ile Thr
                85                  90                  95
Asn Phe Thr Met Leu Ile Gly Asp Lys Val Tyr Gln Gly Glu Ile Thr
            100                 105                 110
Glu Arg Glu Lys Lys Ser Gly Asp Arg Val Lys Glu Lys Arg Asn Lys
        115                 120                 125
Thr Thr Glu Glu Asn Gly Glu Lys Gly Thr Glu Ile Phe Arg Ala Ser
    130                 135                 140
Ala Val Ile Pro Ser Lys Asp Lys Ala Ala Phe Phe Leu Ser Tyr Glu
145                 150                 155                 160
Glu Leu Leu Gln Arg Arg Leu Gly Lys Tyr Glu His Ser Ile Ser Val
                165                 170                 175
Arg Pro Gln Gln Leu Ser Gly Arg Leu Ser Val Asp Val Asn Ile Leu
            180                 185                 190
Glu Ser Ala Gly Ile Ala Ser Leu Glu Val
        195                 200         202
```

<210> 1484
<211> 477
<212> PRT
<213> Homo sapiens

<400> 1484
Met Pro Gln Leu Ser Leu Ser Trp Leu Gly Leu Gly Gln Val Ala Ala
1 5 10 15
Phe Pro Trp Leu Leu Leu Leu Leu Ala Gly Ala Ser Arg Leu Leu Ala
20 25 30
Gly Phe Leu Ala Trp Thr Tyr Ala Phe Tyr Asp Asn Cys Arg Arg Leu
35 40 45
Gln Tyr Phe Pro Gln Pro Pro Lys Gln Lys Trp Phe Trp Gly Gln Pro
50 55 60
Gly Pro Pro Ala Ile Ala Pro Lys Asp Asp Leu Ser Ile Arg Phe Leu
65 70 75 80
Lys Pro Trp Leu Gly Glu Gly Ile Leu Leu Ser Gly Gly Asp Lys Trp
85 90 95
Ser Arg His Arg Arg Met Leu Thr Pro Ala Phe His Phe Asn Ile Leu
100 105 110
Lys Ser Tyr Ile Thr Ile Phe Asn Lys Ser Ala Asn Ile Met Leu Asp
115 120 125
Lys Trp Gln His Leu Ala Ser Glu Gly Ser Ser Cys Leu Asp Met Phe
130 135 140
Glu His Ile Ser Leu Met Thr Leu Asp Ser Leu Gln Lys Cys Ile Phe
145 150 155 160
Ser Phe Asp Ser His Cys Gln Glu Arg Pro Ser Glu Tyr Ile Ala Thr
165 170 175
Ile Leu Glu Leu Ser Ala Leu Val Glu Lys Arg Ser Gln His Ile Leu
180 185 190
Gln His Met Asp Phe Leu Tyr Tyr Leu Ser His Asp Gly Arg Arg Phe
195 200 205
His Arg Ala Cys Arg Leu Val His Asp Phe Thr Asp Ala Val Ile Arg
210 215 220
Glu Arg Arg Arg Thr Leu Pro Thr Gln Gly Ile Asp Asp Phe Phe Lys
225 230 235 240
Asp Lys Ala Lys Ser Lys Thr Leu Asp Phe Ile Asp Val Leu Leu Leu
245 250 255
Ser Lys Asp Glu Asp Gly Lys Ala Leu Ser Asp Glu Asp Ile Arg Ala
260 265 270
Glu Ala Asp Thr Phe Met Phe Gly Gly His Asp Thr Thr Ala Ser Gly
275 280 285
Leu Ser Trp Val Leu Tyr Asn Leu Ala Arg His Pro Glu Tyr Gln Glu
290 295 300
Arg Cys Arg Gln Glu Val Gln Glu Leu Leu Lys Asp Arg Asp Pro Lys
305 310 315 320
Glu Ile Glu Trp Asp Asp Leu Ala Gln Leu Pro Phe Leu Thr Met Cys
325 330 335
Val Lys Glu Ser Leu Arg Leu His Pro Pro Ala Pro Phe Ile Ser Arg
340 345 350
Cys Cys Thr Gln Asp Ile Val Leu Pro Asp Gly Arg Val Ile Pro Lys
355 360 365
Gly Ile Thr Cys Leu Ile Asp Ile Ile Gly Val His His Asn Pro Thr
370 375 380
Val Trp Pro Asp Pro Glu Val Tyr Asp Pro Phe Arg Phe Asp Pro Glu

```
385                 390                 395                 400
Asn Ser Lys Gly Arg Ser Pro Leu Ala Phe Ile Pro Phe Ser Ala Gly
                405                 410                 415
Pro Arg Asn Cys Ile Gly Gln Ala Phe Ala Met Ala Glu Met Lys Val
            420                 425                 430
Val Leu Ala Leu Met Leu Leu His Phe Arg Phe Leu Pro Asp His Thr
        435                 440                 445
Glu Pro Arg Arg Lys Leu Glu Leu Ile Met Arg Ala Glu Gly Gly Leu
    450                 455                 460
Trp Leu Arg Val Glu Pro Leu Asn Val Ser Leu Gln  *
465                 470                 475 476


     <210> 1485
     <211> 67
     <212> PRT
     <213> Homo sapiens


     <400> 1485
Met Ala Cys Cys Leu Phe Leu Asn Gly Ser Trp Leu Ser Met Ala Leu
 1               5                  10                  15
Lys Phe Phe Asn Cys Trp Gly Lys Lys Ile Lys Arg Ile Ile Phe Tyr
            20                  25                  30
Val Lys Ile Met Lys Phe Lys Phe Gln Cys Pro Gln Ile Asn Thr Ala
        35                  40                  45
Thr Tyr Ile His Leu His Gly Cys Phe Cys Thr Ser Met Ala Glu Leu
    50                  55                  60
Ser Ser  *
 65  66


     <210> 1486
     <211> 93
     <212> PRT
     <213> Homo sapiens


     <400> 1486
Met Gly Ser Ser Val Leu Ser Ile Trp Ile Leu Ser Pro Ser Ile Tyr
 1               5                  10                  15
Pro Ile Leu Ser Pro Leu Ala Met Pro Cys Leu Ser Arg Thr Asp Leu
            20                  25                  30
Ile Arg Val Arg Arg Ile Gln Gly Ala Trp Pro Ser Glu Gly Thr Ala
        35                  40                  45
Ser Ser Ile Arg Gly Trp Val Leu Thr Lys Leu Arg Met Ser Ser Gly
    50                  55                  60
Lys Ala Leu Glu Ala Leu Tyr Cys Ile Pro Gly Ala Ala Gln His Pro
 65                 70                  75                  80
Gly Leu Gly Val Thr Arg Val Trp Ser Gly Arg Thr  *
                85                  90      92


     <210> 1487
     <211> 88
     <212> PRT
```

<213> Homo sapiens

<400> 1487

```
Met Gln Lys Val Thr Leu Gly Leu Leu Val Phe Leu Ala Gly Phe Pro
 1               5                  10                  15
Val Leu Asp Ala Asn Asp Leu Glu Asp Lys Asn Ser Pro Phe Tyr Tyr
            20                  25                  30
Asp Trp His Ser Leu Gln Val Gly Gly Leu Ile Cys Ala Gly Val Leu
        35                  40                  45
Cys Ala Met Gly Ile Ile Ile Val Met Ser Ala Lys Cys Lys Cys Lys
    50                  55                  60
Phe Gly Gln Lys Ser Gly His His Pro Gly Glu Thr Pro Pro Leu Ile
65                  70                  75                  80
Thr Pro Gly Ser Ala Gln Ser  *
                85      87
```

<210> 1488
<211> 268
<212> PRT
<213> Homo sapiens

<400> 1488

```
Met Gly Ser Ala Cys Ile Lys Val Thr Lys Tyr Phe Leu Phe Leu Phe
 1               5                  10                  15
Asn Leu Ile Phe Phe Ile Leu Gly Ala Val Ile Leu Gly Phe Gly Val
            20                  25                  30
Trp Ile Leu Ala Asp Lys Ser Ser Phe Ile Ser Val Leu Gln Thr Ser
        35                  40                  45
Ser Ser Ser Leu Arg Met Gly Ala Tyr Val Phe Ile Gly Val Gly Ala
    50                  55                  60
Val Thr Met Leu Met Gly Phe Leu Gly Cys Ile Gly Ala Val Asn Glu
65                  70                  75                  80
Val Arg Cys Leu Leu Gly Leu Tyr Phe Ala Phe Leu Leu Leu Ile Leu
                85                  90                  95
Ile Ala Gln Val Thr Ala Gly Ala Leu Phe Tyr Phe Asn Met Gly Lys
            100                 105                 110
Leu Lys Gln Glu Met Gly Gly Ile Val Thr Glu Leu Ile Arg Asp Tyr
        115                 120                 125
Asn Ser Ser Arg Glu Asp Ser Leu Gln Asp Ala Trp Asp Tyr Val Gln
    130                 135                 140
Ala Gln Val Lys Cys Cys Gly Trp Val Ser Phe Tyr Asn Trp Thr Asp
145                 150                 155                 160
Asn Ala Glu Leu Met Asn Arg Pro Glu Val Thr Tyr Pro Cys Ser Cys
                165                 170                 175
Glu Val Lys Gly Glu Glu Asp Asn Ser Leu Ser Val Arg Lys Gly Phe
            180                 185                 190
Cys Glu Ala Pro Gly Asn Arg Thr Gln Ser Gly Asn His Pro Glu Asp
        195                 200                 205
Trp Pro Val Tyr Gln Glu Gly Cys Met Glu Lys Val Gln Ala Trp Leu
    210                 215                 220
Gln Glu Asn Leu Gly Ile Ile Leu Gly Val Gly Val Gly Val Ala Ile
225                 230                 235                 240
Ile Glu Leu Leu Gly Met Val Leu Ser Ile Cys Leu Cys Arg His Val
                245                 250                 255
His Ser Glu Asp Tyr Ser Lys Val Pro Lys Tyr  *
```

260 265 267

<210> 1489
<211> 832
<212> PRT
<213> Homo sapiens

<400> 1489
Met Thr Leu Ala Leu Ala Tyr Leu Leu Ala Leu Pro Gln Val Leu Asp
1 5 10 15
Ala Asn Arg Cys Phe Glu Lys Gln Ser Pro Ser Ala Leu Ser Leu Gln
20 25 30
Leu Ala Ala Tyr Tyr Tyr Ser Leu Gln Ile Tyr Ala Arg Leu Ala Pro
35 40 45
Cys Phe Arg Asp Lys Cys His Pro Leu Tyr Arg Ala Asp Pro Lys Glu
50 55 60
Leu Ile Lys Met Val Thr Arg His Val Thr Arg His Glu His Glu Ala
65 70 75 80
Trp Pro Glu Asp Leu Ile Ser Leu Thr Lys Gln Leu His Cys Tyr Asn
85 90 95
Glu Arg Leu Leu Asp Phe Thr Gln Ala Gln Ile Leu Gln Gly Leu Arg
100 105 110
Lys Gly Val Asp Val Gln Arg Phe Thr Ala Asp Asp Gln Tyr Lys Arg
115 120 125
Glu Thr Ile Leu Gly Leu Ala Glu Thr Leu Glu Glu Ser Val Tyr Ser
130 135 140
Ile Ala Ile Ser Leu Ala Gln Arg Tyr Ser Val Ser Arg Trp Glu Val
145 150 155 160
Phe Met Thr His Leu Glu Phe Leu Phe Thr Asp Ser Gly Leu Ser Thr
165 170 175
Leu Glu Ile Glu Asn Arg Ala Gln Asp Leu His Leu Phe Glu Thr Leu
180 185 190
Lys Thr Asp Pro Glu Ala Phe His Gln His Met Val Lys Tyr Ile Tyr
195 200 205
Pro Thr Ile Gly Gly Phe Asp His Glu Arg Leu Gln Tyr Tyr Phe Thr
210 215 220
Leu Leu Glu Asn Cys Gly Cys Ala Asp Leu Gly Asn Cys Ala Ile Lys
225 230 235 240
Pro Glu Thr His Ile Arg Leu Leu Lys Lys Phe Lys Val Val Ala Ser
245 250 255
Gly Leu Asn Tyr Lys Lys Leu Thr Asp Glu Asn Met Ser Pro Leu Glu
260 265 270
Ala Leu Glu Pro Val Leu Ser Ser Gln Asn Ile Leu Ser Ile Ser Lys
275 280 285
Leu Val Pro Lys Ile Pro Glu Lys Asp Gly Gln Met Leu Ser Pro Ser
290 295 300
Ser Leu Tyr Thr Ile Trp Leu Gln Lys Leu Phe Trp Thr Gly Asp Pro
305 310 315 320
His Leu Ile Lys Gln Val Pro Gly Ser Ser Pro Glu Trp Leu His Ala
325 330 335
Tyr Asp Val Cys Met Lys Tyr Phe Asp Arg Leu His Pro Gly Asp Leu
340 345 350
Ile Thr Val Val Asp Ala Val Thr Phe Ser Pro Lys Ala Val Thr Lys
355 360 365
Leu Ser Val Glu Ala Arg Lys Glu Met Thr Arg Lys Ala Ile Lys Thr
370 375 380

```
Val Lys His Phe Ile Glu Lys Pro Arg Lys Arg Asn Ser Glu Asp Glu
385                 390                 395                 400
Ala Gln Glu Ala Lys Asp Ser Lys Val Thr Tyr Ala Asp Thr Leu Asn
                405                 410                 415
His Leu Glu Lys Ser Leu Ala His Leu Glu Thr Leu Ser His Ser Phe
            420                 425                 430
Ile Leu Ser Leu Lys Asn Ser Glu Gln Glu Thr Leu Gln Lys Tyr Ser
        435                 440                 445
His Leu Tyr Asp Leu Ser Arg Ser Glu Lys Glu Lys Leu His Asp Glu
    450                 455                 460
Ala Val Ala Ile Cys Leu Asp Gly Gln Pro Leu Ala Met Ile Gln Gln
465                 470                 475                 480
Leu Leu Glu Val Ala Val Gly Pro Leu Asp Ile Ser Pro Lys Asp Ile
                485                 490                 495
Val Gln Ser Ala Ile Met Lys Ile Ile Ser Ala Leu Ser Gly Gly Ser
            500                 505                 510
Ala Asp Leu Gly Gly Pro Arg Asp Pro Leu Lys Val Leu Glu Gly Val
        515                 520                 525
Val Ala Ala Val His Ala Ser Val Asp Lys Gly Glu Glu Leu Val Ser
    530                 535                 540
Pro Glu Asp Leu Leu Glu Trp Leu Arg Pro Phe Cys Ala Asp Asp Ala
545                 550                 555                 560
Trp Pro Val Arg Pro Arg Ile His Val Leu Gln Ile Leu Gly Gln Ser
                565                 570                 575
Phe His Leu Thr Glu Glu Asp Ser Lys Leu Leu Val Phe Phe Arg Thr
            580                 585                 590
Glu Ala Ile Leu Lys Ala Ser Trp Pro Gln Arg Gln Val Asp Ile Ala
        595                 600                 605
Asp Ile Glu Asn Glu Glu Asn Arg Tyr Cys Leu Phe Met Glu Leu Leu
    610                 615                 620
Glu Ser Ser His His Glu Ala Glu Phe Gln His Leu Val Leu Leu Leu
625                 630                 635                 640
Gln Ala Trp Pro Pro Met Lys Ser Glu Tyr Val Ile Thr Asn Asn Pro
                645                 650                 655
Trp Val Arg Leu Ala Thr Val Met Leu Thr Arg Cys Thr Met Glu Asn
            660                 665                 670
Lys Glu Gly Leu Gly Asn Glu Val Leu Lys Met Cys Arg Ser Leu Tyr
        675                 680                 685
Asn Thr Lys Gln Met Leu Pro Ala Glu Gly Val Lys Glu Leu Cys Leu
    690                 695                 700
Leu Leu Leu Asn Gln Ser Leu Leu Leu Pro Ser Leu Lys Leu Leu Leu
705                 710                 715                 720
Glu Ser Arg Asp Glu His Leu His Glu Met Ala Leu Glu Gln Ile Thr
                725                 730                 735
Ala Val Thr Thr Val Asn Asp Ser Asn Cys Asp Gln Glu Leu Leu Ser
            740                 745                 750
Leu Leu Leu Asp Ala Lys Leu Leu Val Lys Cys Val Ser Thr Pro Phe
        755                 760                 765
Tyr Pro Arg Ile Val Asp His Leu Leu Ala Ser Leu Gln Gln Gly Arg
    770                 775                 780
Trp Asp Ala Glu Glu Leu Gly Arg His Leu Arg Glu Ala Gly His Glu
785                 790                 795                 800
Ala Glu Ala Gly Ser Leu Leu Leu Ala Val Arg Gly Thr His Gln Ala
                805                 810                 815
Phe Arg Thr Phe Ser Thr Ala Leu Arg Ala Ala Gln His Trp Val  *
            820                 825                 830 831
```

<210> 1490
<211> 55
<212> PRT
<213> Homo sapiens

<400> 1490
Met Trp Phe Leu Leu Val Ser Val Val Cys Leu Tyr Gly Ile Gly Glu
1 5 10 15
Gly Asn Phe Phe Ser Leu Ala Ser Val Phe Ser Leu Leu Ser Leu Cys
20 25 30
Leu His Leu Leu Leu Trp Lys Arg Ala Phe Asp Arg Thr Asp Val Leu
35 40 45
Thr Ser Glu Trp Ile Phe *
50 54

<210> 1491
<211> 134
<212> PRT
<213> Homo sapiens

<400> 1491
Met Thr Thr Thr Phe Pro Pro Arg Lys Met Val Ala Gln Phe Leu Leu
1 5 10 15
Val Ala Gly Asn Val Ala Asn Ile Thr Thr Val Ser Leu Trp Glu Glu
20 25 30
Phe Ser Ser Ser Asp Leu Ala Asp Leu Arg Phe Leu Asp Met Ser Gln
35 40 45
Asn Gln Phe Gln Tyr Leu Pro Asp Gly Phe Leu Arg Lys Met Pro Ser
50 55 60
Leu Ser His Leu Asn Leu His Gln Asn Cys Leu Met Thr Leu His Ile
65 70 75 80
Arg Glu His Glu Pro Pro Gly Ala Leu Thr Glu Leu Asp Leu Ser His
85 90 95
Asn Gln Leu Ser Glu Leu His Leu Ala Pro Gly Leu Ala Ser Cys Leu
100 105 110
Gly Ser Leu Arg Leu Phe Asn Leu Ser Ser Asn Gln Leu Leu Gly Val
115 120 125
Pro Pro Gly Pro Leu Tyr
130 134

<210> 1492
<211> 71
<212> PRT
<213> Homo sapiens

<400> 1492
Met Arg Ser Glu Trp Phe Tyr Lys Trp Phe Phe Pro Pro Phe Ala Leu
1 5 10 15
His Phe Ser Leu Leu Pro Pro Cys Glu Glu Gly His Val Cys Leu Pro
20 25 30
Met Cys His Glu Cys Lys Phe Pro Glu Ala Ser Pro Ala Thr Met Asn
35 40 45

```
Cys Glu Ser Ile Lys Pro Leu Phe Leu Ile Asn Tyr Pro Val Ser Asn
     50                  55                  60
Lys Ser Leu Leu Ala Thr  *
 65                  70
```

<210> 1493
<211> 78
<212> PRT
<213> Homo sapiens

<400> 1493

```
Met Trp Ile Tyr Phe Trp Thr Leu Asn Ser Val Pro Val Ile Tyr Met
  1               5                  10                  15
Ser Thr Leu Met Ser Ile Pro His Tyr Phe Asp Tyr Cys Cys Phe Ile
             20                  25                  30
Val Ser Asp Ile Met Leu Pro Glu Ile Thr Phe Ser Thr Phe Ile Leu
         35                  40                  45
Leu Leu Met Val Ala Leu Ala Ile Arg Gly Pro Leu His Phe Arg Arg
     50                  55                  60
His Phe Arg Ile Asn Leu Ser Ile Ala Thr Lys Asn Ala  *
 65                  70                  75      77
```

<210> 1494
<211> 121
<212> PRT
<213> Homo sapiens

<400> 1494

```
Met Ala Gly Leu Asn Cys Gly Val Ser Ile Ala Leu Leu Gly Val Leu
  1               5                  10                  15
Leu Leu Gly Ala Ala Arg Leu Pro Arg Gly Ala Glu Ala Phe Glu Ile
             20                  25                  30
Ala Leu Pro Arg Glu Ser Asn Ile Thr Val Leu Ile Lys Leu Gly Thr
         35                  40                  45
Pro Thr Leu Leu Ala Lys Pro Cys Tyr Ile Val Ile Ser Lys Arg His
     50                  55                  60
Ile Thr Met Leu Ser Ile Lys Ser Gly Glu Arg Ile Val Phe Thr Phe
 65                  70                  75                  80
Ser Cys Gln Ser Pro Glu Asn His Phe Val Ile Glu Ile Gln Lys Asn
                 85                  90                  95
Ile Asp Cys Met Ser Gly Pro Cys Pro Phe Gly Glu Val Gln Leu Gln
            100                 105                 110
Pro Ser Thr Ser Leu Leu Pro Thr Leu
        115                 120 121
```

<210> 1495
<211> 91
<212> PRT
<213> Homo sapiens

```
     <400> 1495
Met Glu Asn Cys Val Gly Glu Arg Thr His Pro Leu Phe Val Val Tyr
 1               5                  10                  15
Leu Ala Leu Gln Leu Val Val Leu Leu Trp Gly Leu Tyr Leu Ala Trp
            20                  25                  30
Ser Gly Leu Arg Phe Phe Gln Pro Trp Gly Leu Trp Leu Arg Ser Ser
        35                  40                  45
Gly Leu Leu Phe Ala Thr Phe Gln Leu Leu Ser Leu Phe Ser Leu Val
    50                  55                  60
Ala Ser Leu Leu Leu Val Ser His Leu Tyr Leu Val Ala Ser Asn Thr
65                  70                  75                  80
Thr Thr Trp Glu Phe Ile Ser Ser His His Val
                85                  90  91


     <210> 1496
     <211> 72
     <212> PRT
     <213> Homo sapiens


     <400> 1496
Met Ile Glu Thr Trp Leu Trp Leu Leu Leu Leu Asn Val Gly Gly Thr
 1               5                  10                  15
Gly Gln Trp Ser Gly Pro Thr Phe Arg Arg Glu Asn Val Leu Pro Ala
            20                  25                  30
Ala His Ile Gly Pro Lys Tyr Gly Pro Leu Leu Pro Ser Thr Ala Lys
        35                  40                  45
Gly Thr Val Lys Val Ser Cys Pro Ser Ser Thr Pro His Pro Pro Leu
    50                  55                  60
Gln Gly Lys Gly Thr Pro Asp  *
65                  70  71


     <210> 1497
     <211> 196
     <212> PRT
     <213> Homo sapiens


     <400> 1497
Met Ala Pro Arg Ala Leu Pro Gly Ser Ala Val Leu Ala Ala Ala Val
 1               5                  10                  15
Phe Val Gly Gly Ala Val Ser Ser Pro Leu Val Ala Pro Asp Asn Gly
            20                  25                  30
Ser Ser Arg Thr Leu His Ser Arg Thr Glu Thr Thr Pro Ser Pro Ser
        35                  40                  45
Asn Asp Thr Gly Asn Gly His Pro Glu Tyr Ile Ala Tyr Ala Leu Val
    50                  55                  60
Pro Val Phe Phe Ile Met Gly Leu Phe Gly Val Leu Ile Cys His Leu
65                  70                  75                  80
Leu Lys Lys Lys Gly Tyr Arg Cys Thr Thr Glu Ala Glu Gln Asp Ile
                85                  90                  95
Glu Glu Glu Lys Val Glu Lys Ile Glu Leu Asn Asp Ser Val Asn Glu
            100                 105                 110
Asn Ser Asp Thr Val Gly Gln Ile Val His Tyr Ile Met Lys Asn Glu
        115                 120                 125
```

Ala Asn Ala Asp Val Leu Lys Ala Met Val Ala Asp Asn Ser Leu Tyr
130 135 140
Asp Pro Glu Ser Pro Val Thr Pro Ser Thr Pro Gly Glu Pro Ala Ser
145 150 155 160
Glu Ser Trp Ala Phe Val Thr Arg Gly Asp Ala Arg Glu Ala Arg Leu
165 170 175
Trp Pro Ser Ser Ala Tyr Gly Gly Arg Cys Cys Arg Glu Gly Cys Val
180 185 190
Ser Ser Val *
195

<210> 1498
<211> 75
<212> PRT
<213> Homo sapiens

<400> 1498
Met Trp Ser Gln Ile Ala Phe Val Arg Ile Pro Phe Cys Phe Ser Leu
1 5 10 15
Leu Ser His Ser Asn Ala Trp Phe Val Gln Lys Ala Ala Ser Gln Arg
20 25 30
Gln Ala Ser Ile Ser Thr Ala Cys His Cys Pro Ala Glu Ala Gly Gly
35 40 45
Glu Arg Ile Thr Val Ser Thr Thr Gly Ala Gln Arg Asn Ala Ala Met
50 55 60
Val Pro Asp Leu Gln Ser Pro Arg Arg Ser *
65 70 74

<210> 1499
<211> 62
<212> PRT
<213> Homo sapiens

<400> 1499
Met Pro Ser Leu Met Met Val Leu Glu Ala Arg Phe Val Ser Ser Cys
1 5 10 15
Leu Ile Phe Pro Ser Arg Ala Met Pro Leu Leu Ser Arg Leu Leu Ala
20 25 30
Ser Lys Gly Ser Ser Val Asn Val Leu Val Lys Val Leu Phe Gly Gly
35 40 45
Thr Phe Ser Cys Ala Ser Ser Ile Ala Thr Gly Leu Thr *
50 55 60 61

<210> 1500
<211> 138
<212> PRT
<213> Homo sapiens

<400> 1500
Met Pro Ile Trp Lys Pro Phe Met Ala Trp Met Ala Ala Trp Ala Leu

```
 1               5                   10                  15
Ala Val Leu Ser Lys Leu Thr Lys Pro Ile His Leu Leu Trp Met Val
            20                  25                  30
Ala Arg Ser Ile Asn Thr Leu Glu Glu Met Ile Leu Pro Lys Gly Thr
        35                  40                  45
Asn Ile Cys Val Ser Ser Val Ser Pro Asn Ser Phe Ser Leu Leu Leu
    50                  55                  60
Leu Gln Glu Gly Arg Arg Leu Glu Asp Ala Val Arg Asp Gly Arg Asp
65                  70                  75                  80
Gly Arg Gly Gly Ala His Gly Cys Val Leu Leu Asp Ser Gly Glu Gly
                85                  90                  95
Arg Met Gln Cys Leu Gly His Ser Arg Ala Leu Ser Trp Val Trp His
            100                 105                 110
Lys Ala Ile Gly Ile Asp Glu Phe Pro Gly Gln Gly Ala His Leu Glu
        115                 120                 125
Arg Ala Arg His Leu Pro Ser His Trp  *
    130                 135     137
```

```
<210> 1501
<211> 82
<212> PRT
<213> Homo sapiens


<400> 1501
Met Ile Leu Phe Thr Arg Ala Trp Phe Glu Leu Val Thr Leu Val Gln
 1               5                   10                  15
Phe Ile Ile Gly Ser Gln Met Leu Tyr Pro Tyr Leu His Ile Glu Glu
            20                  25                  30
Phe Val Ile Arg Lys Leu Pro Val Leu Leu Tyr Arg Lys Ser Val Ile
        35                  40                  45
Arg Tyr Gln Met Ala Ser Ser Pro Cys Leu Gln Met Phe Lys Gln Tyr
    50                  55                  60
Cys Gly Trp Ser Arg Lys Ser Leu Arg His Ala Val Lys Cys Arg Ala
65                  70                  75                  80
Arg  *
 81
```

```
<210> 1502
<211> 54
<212> PRT
<213> Homo sapiens


<400> 1502
Met Leu Leu Phe Leu Gly Phe Phe Ile Cys Ser Leu Phe Phe Ser Glu
 1               5                   10                  15
Leu Ser Thr Gly Thr Thr His Ser Leu Glu Ser Tyr Gln Ile Leu Leu
            20                  25                  30
Ser Lys Phe Phe Arg His Pro Leu Cys Thr Arg Thr Phe Arg Ile Leu
        35                  40                  45
Pro Pro Phe His Phe  *
    50          53
```

<210> 1503
<211> 62
<212> PRT
<213> Homo sapiens

<400> 1503
Met Gly Trp Pro Pro Ser Leu Trp Val Leu Ala Leu Ala Tyr Cys Cys
1 5 10 15
Lys Ala Pro Gln Arg Leu Cys Ser Gly Ser Ser Pro Cys Arg Phe Ser
20 25 30
Ser Arg Met Ser Ala Ser Pro Ala Thr Asn Arg Asn Glu Asn Thr Thr
35 40 45
Ser Trp Ile Ala Ser Leu His Lys Tyr Val Ile Ser Gln *
50 55 60 61

<210> 1504
<211> 46
<212> PRT
<213> Homo sapiens

<400> 1504
Met Trp Lys Gln Ile Ser Pro Ile Gly Asn Leu Val Thr Ala Ile Phe
1 5 10 15
Phe Cys Val Leu Cys Gln Gln Arg Tyr Gln Trp Leu Ala Arg Asp Ala
20 25 30
Phe Asn Thr Gln Ser Ile Leu Ser Pro Pro Ile Trp Val *
35 40 45

<210> 1505
<211> 48
<212> PRT
<213> Homo sapiens

<400> 1505
Met Val Ala Val Ser Leu Leu Cys Pro Trp Pro Ser Ser Trp Asn Arg
1 5 10 15
Arg Ser Cys Gly Arg Ser His Arg Asn Leu Gly Leu Phe Thr Ala Phe
20 25 30
Leu Ser Val Pro Glu Phe Val Ile Phe Gly Ala Cys Arg Tyr Trp *
35 40 45 47

<210> 1506
<211> 190
<212> PRT
<213> Homo sapiens

<400> 1506
Met Trp Leu Leu Gly Pro Leu Cys Leu Leu Leu Ser Ser Ala Ala Glu

```
 1               5                   10                  15
Ser Gln Leu Leu Pro Gly Asn Asn Phe Thr Asn Glu Cys Asn Ile Pro
            20                  25                  30
Gly Asn Phe Val Cys Ser Asn Gly Arg Cys Ile Pro Gly Ala Trp Gln
        35                  40                  45
Cys Asp Gly Leu Pro Asp Cys Phe Asp Lys Ser Asp Glu Lys Glu Cys
    50                  55                  60
Pro Lys Ala Lys Ser Lys Cys Gly Pro Thr Phe Phe Pro Cys Ala Ser
65                  70                  75                  80
Gly Ile His Cys Ile Ile Gly Arg Phe Arg Cys Asn Gly Phe Glu Asp
                85                  90                  95
Cys Pro Asp Gly Ser Asp Glu Glu Asn Cys Thr Ala Asn Pro Leu Leu
            100                 105                 110
Cys Ser Thr Ala Arg Tyr His Cys Lys Asn Gly Leu Cys Ile Asp Lys
        115                 120                 125
Ser Phe Ile Cys Asp Gly Gln Asn Asn Cys Gln Asp Asn Ser Asp Glu
    130                 135                 140
Glu Ser Cys Glu Ser Ser Gln Val Phe Arg Pro Gln Val Ser Glu Trp
145                 150                 155                 160
Gln Ala Arg Pro Arg Asp Leu Cys Ala Arg Trp Asn Ile Pro Phe Leu
                165                 170                 175
Gly Arg Leu Glu Arg Pro Trp Ser Phe Thr Ser Ser Gln Gln
            180                 185                 190
```

```
     <210> 1507
     <211> 60
     <212> PRT
     <213> Homo sapiens
```

```
     <400> 1507
Met Tyr Arg Pro Ala Pro Pro Arg Gln Asn Arg Gln Leu His Pro Tyr
 1               5                   10                  15
Leu Leu Ala Ser Trp Pro Lys Ala Leu Asn Cys Thr Leu Cys Val Cys
            20                  25                  30
Val Cys Val Cys Ala Arg Val Cys Ala Cys Val Cys Met Trp Ser Val
        35                  40                  45
Thr Ser Leu Trp Leu Thr Cys Leu Ser Gly Val  *
    50                  55              59
```

```
     <210> 1508
     <211> 48
     <212> PRT
     <213> Homo sapiens
```

```
     <400> 1508
Met Ser His His Cys Ala Trp Pro Lys Asn Phe Leu Leu Lys Met Leu
 1               5                   10                  15
Ser Thr Gly Arg Val Gln Trp Leu Met Pro Ile Ile Phe Leu Phe Phe
            20                  25                  30
Gln Lys Met Gly Gly Asn Met Val Gly Ser Gln Leu Lys Leu Ser  *
        35                  40                  45      47
```

<210> 1509
<211> 85
<212> PRT
<213> Homo sapiens

<400> 1509
Met Thr Gly Ser Arg Cys Glu Glu His Val Phe Ser Gln Gln Gln Pro
1 5 10 15
Gly His Ile Ala Ser Ile Leu Ile Pro Leu Leu Leu Leu Leu Leu Leu
20 25 30
Val Leu Ala Ala Gly Val Val Phe Trp Tyr Lys Arg Arg Val Gln Gly
35 40 45
Ala Lys Gly Phe His His Gln Arg Met Thr Asn Gly Ala Met Asn Val
50 55 60
Glu Ile Gly Asn Pro Thr Tyr Lys Met Tyr Glu Gly Gly Glu Pro Asp
65 70 75 80
Asp Val Gly Gly Leu
85

<210> 1510
<211> 55
<212> PRT
<213> Homo sapiens

<400> 1510
Met Ala Ile Ser Trp Lys Pro Thr Gly Leu Pro Trp His Ser Met Leu
1 5 10 15
Gln Val Leu Leu Ala Ala Trp Leu Pro Gly Pro Thr Pro Thr Pro His
20 25 30
Ser Ala Leu Pro Ser Phe Ser Pro Pro Pro Ser Leu Pro Pro Lys Met
35 40 45
Cys Leu Pro Lys Cys Cys *
50 54

<210> 1511
<211> 108
<212> PRT
<213> Homo sapiens

<400> 1511
Met Val Gly Phe Gly Ala Asn Arg Arg Ala Gly Arg Leu Pro Ser Leu
1 5 10 15
Val Leu Gly Val Leu Leu Val Val Ile Val Val Leu Ala Phe Asn Tyr
20 25 30
Trp Ser Ile Ser Ser Arg His Val Leu Leu Gln Glu Glu Val Ala Glu
35 40 45
Leu Gln Gly Gln Val Gln Arg Thr Glu Val Ala Arg Gly Arg Leu Glu
50 55 60
Lys Arg Asn Ser Asp Leu Phe Ala Val Val Gly His Ala Gln Glu Thr
65 70 75 80
Asp Arg Pro Glu Gly Gly Arg Leu Arg Pro Pro Gln Gln Pro Ala Ala

```
                85                  90                  95
Gly Gln Arg Gly Pro Arg Glu Glu Met Arg Gly  *
            100                 105     107
```

<210> 1512
<211> 119
<212> PRT
<213> Homo sapiens

<400> 1512

```
Met Val Ala Arg Val Trp Ser Leu Met Arg Phe Leu Ile Lys Gly Ser
 1               5                  10                  15
Val Ala Gly Gly Ala Val Tyr Leu Val Tyr Asp Gln Glu Leu Leu Gly
            20                  25                  30
Pro Ser Asp Lys Ser Gln Ala Ala Leu Gln Lys Ala Gly Glu Val Val
        35                  40                  45
Pro Pro Ala Met Tyr Gln Phe Ser Gln Tyr Val Cys Gln Gln Thr Gly
    50                  55                  60
Leu Gln Ile Pro Gln Leu Pro Ala Pro Pro Lys Ile Tyr Phe Pro Ile
65                  70                  75                  80
Arg Asp Ser Trp Asn Ala Gly Ile Met Thr Val Met Ser Ala Leu Ser
                85                  90                  95
Val Ala Pro Ser Lys Ala Arg Glu Tyr Ser Lys Glu Gly Trp Glu Tyr
            100                 105                 110
Val Lys Ala Arg Thr Lys  *
        115         118
```

<210> 1513
<211> 973
<212> PRT
<213> Homo sapiens

<400> 1513

```
Met Val Lys Ser Lys Trp Gly Leu Ala Leu Ala Ala Val Val Thr Val
 1               5                  10                  15
Leu Ser Ser Leu Leu Met Ser Val Gly Leu Cys Thr Leu Phe Gly Leu
            20                  25                  30
Thr Pro Thr Leu Asn Gly Gly Glu Ile Phe Pro Tyr Leu Val Val Val
        35                  40                  45
Ile Gly Leu Glu Asn Val Leu Val Leu Thr Lys Ser Val Val Ser Thr
    50                  55                  60
Pro Val Asp Leu Glu Val Lys Leu Arg Ile Ala Gln Gly Leu Ser Ser
65                  70                  75                  80
Glu Ser Trp Ser Ile Met Lys Asn Met Ala Thr Glu Leu Gly Ile Ile
                85                  90                  95
Leu Ile Gly Tyr Phe Thr Leu Val Pro Ala Ile Gln Glu Phe Cys Leu
            100                 105                 110
Phe Ala Val Val Gly Leu Val Ser Asp Phe Phe Leu Gln Met Leu Phe
        115                 120                 125
Phe Thr Thr Val Leu Ser Ile Asp Ile Arg Arg Met Glu Leu Ala Asp
    130                 135                 140
Leu Asn Lys Arg Leu Pro Pro Glu Ala Cys Leu Pro Ser Ala Lys Pro
145                 150                 155                 160
```

```
Val Gly Gln Pro Thr Arg Tyr Glu Arg Gln Leu Ala Val Arg Pro Ser
                165                 170                 175
Thr Pro His Thr Ile Thr Leu Gln Pro Ser Ser Phe Arg Asn Leu Arg
            180                 185                 190
Leu Pro Lys Arg Leu Arg Val Val Tyr Phe Leu Ala Arg Thr Arg Leu
        195                 200                 205
Ala Gln Arg Leu Ile Met Ala Gly Thr Val Val Trp Ile Gly Ile Leu
    210                 215                 220
Val Tyr Thr Asp Pro Ala Gly Leu Arg Asn Tyr Leu Ala Ala Gln Val
225                 230                 235                 240
Thr Glu Gln Ser Pro Leu Gly Glu Gly Ala Leu Ala Pro Met Pro Val
                245                 250                 255
Pro Ser Gly Met Leu Pro Pro Ser His Pro Asp Pro Ala Phe Ser Ile
            260                 265                 270
Phe Pro Pro Asp Ala Pro Lys Leu Pro Glu Asn Gln Thr Ser Pro Gly
        275                 280                 285
Glu Ser Pro Glu Arg Gly Gly Pro Ala Glu Val Val His Asp Ser Pro
    290                 295                 300
Val Pro Glu Val Thr Trp Gly Pro Glu Asp Glu Glu Leu Trp Arg Lys
305                 310                 315                 320
Leu Ser Phe Arg His Trp Pro Thr Leu Phe Ser Tyr Tyr Asn Ile Thr
                325                 330                 335
Leu Ala Lys Arg Tyr Ile Ser Leu Leu Pro Val Ile Pro Val Thr Leu
            340                 345                 350
Arg Leu Asn Pro Arg Glu Ala Leu Glu Gly Arg His Pro Gln Asp Gly
        355                 360                 365
Arg Ser Ala Trp Pro Pro Pro Gly Pro Ile Pro Ala Gly His Trp Glu
    370                 375                 380
Ala Gly Pro Lys Gly Pro Gly Gly Val Gln Ala His Gly Asp Val Thr
385                 390                 395                 400
Leu Tyr Lys Val Ala Ala Leu Gly Leu Ala Thr Gly Ile Val Leu Val
                405                 410                 415
Leu Leu Leu Leu Cys Leu Tyr Arg Val Leu Cys Pro Arg Asn Tyr Gly
            420                 425                 430
Gln Leu Gly Gly Gly Pro Gly Arg Arg Arg Arg Gly Glu Leu Pro Cys
        435                 440                 445
Asp Asp Tyr Gly Tyr Ala Pro Pro Glu Thr Glu Ile Val Pro Leu Val
    450                 455                 460
Leu Arg Gly His Leu Met Asp Ile Glu Cys Leu Ala Ser Asp Gly Met
465                 470                 475                 480
Leu Leu Val Ser Cys Cys Leu Ala Gly His Val Cys Val Trp Asp Ala
                485                 490                 495
Gln Thr Gly Asp Cys Leu Thr Arg Ile Pro Arg Pro Gly Arg Gln Arg
            500                 505                 510
Arg Asp Ser Gly Val Gly Ser Gly Leu Glu Ala Gln Glu Ser Trp Glu
        515                 520                 525
Arg Leu Ser Asp Gly Gly Lys Ala Gly Pro Glu Glu Pro Gly Asp Ser
    530                 535                 540
Pro Pro Leu Arg His Arg Pro Arg Gly Pro Pro Pro Pro Ser Leu Phe
545                 550                 555                 560
Gly Asp Gln Pro Asp Leu Thr Cys Leu Ile Asp Thr Asn Phe Ser Ala
                565                 570                 575
Gln Pro Arg Ser Ser Gln Pro Thr Gln Pro Glu Pro Arg His Arg Ala
            580                 585                 590
Val Cys Gly Arg Ser Arg Asp Ser Pro Gly Tyr Asp Phe Ser Cys Leu
        595                 600                 605
Val Gln Arg Val Tyr Gln Glu Glu Gly Leu Ala Ala Val Cys Thr Pro
    610                 615                 620
Ala Leu Arg Pro Pro Ser Pro Gly Pro Val Leu Ser Gln Ala Pro Glu
```

625 630 635 640
Asp Glu Gly Gly Ser Pro Glu Lys Gly Ser Pro Ser Leu Ala Trp Ala
645 650 655
Pro Ser Ala Glu Gly Ser Ile Trp Ser Leu Glu Leu Gln Gly Asn Leu
660 665 670
Ile Val Val Gly Arg Ser Ser Gly Arg Leu Glu Val Trp Asp Ala Ile
675 680 685
Glu Gly Val Leu Cys Cys Ser Ser Glu Glu Val Ser Ser Gly Ile Thr
690 695 700
Ala Leu Val Phe Leu Asp Lys Arg Ile Val Ala Ala Arg Leu Asn Gly
705 710 715 720
Ser Leu Asp Phe Phe Ser Leu Glu Thr His Thr Ala Leu Ser Pro Leu
725 730 735
Gln Phe Arg Gly Thr Pro Gly Arg Gly Ser Ser Pro Ala Ser Pro Val
740 745 750
Tyr Ser Ser Ser Asp Thr Val Ala Cys His Leu Thr His Thr Val Pro
755 760 765
Cys Ala His Gln Lys Pro Ile Thr Ala Leu Lys Ala Ala Ala Gly Arg
770 775 780
Leu Val Thr Gly Ser Gln Asp His Thr Leu Arg Val Phe Arg Leu Glu
785 790 795 800
Asp Ser Cys Cys Leu Phe Thr Leu Gln Gly His Ser Gly Ala Ile Thr
805 810 815
Thr Val Tyr Ile Asp Gln Thr Met Val Leu Ala Ser Gly Gly Gln Asp
820 825 830
Gly Ala Ile Cys Leu Trp Asp Val Leu Thr Gly Ser Arg Val Ser His
835 840 845
Val Phe Ala His Arg Gly Asp Val Thr Ser Leu Thr Cys Thr Thr Ser
850 855 860
Cys Val Ile Ser Ser Gly Leu Asp Asp Leu Ile Ser Ile Trp Asp Arg
865 870 875 880
Ser Thr Gly Ile Lys Phe Tyr Ser Ile Gln Gln Asp Leu Gly Cys Gly
885 890 895
Ala Ser Leu Gly Val Ile Ser Asp Asn Leu Leu Val Thr Gly Gly Gln
900 905 910
Gly Cys Val Ser Phe Trp Asp Leu Asn Tyr Gly Asp Leu Leu Gln Thr
915 920 925
Val Tyr Leu Gly Lys Asn Ser Glu Ala Gln Pro Ala Arg Gln Ile Leu
930 935 940
Val Leu Asp Asn Ala Ala Ile Val Cys Asn Phe Gly Ser Glu Leu Ser
945 950 955 960
Leu Val Tyr Val Pro Ser Val Leu Glu Lys Leu Asp *
965 970 972

<210> 1514
<211> 77
<212> PRT
<213> Homo sapiens

<400> 1514
Met Ile Ser Ser Trp Pro Phe Ser Arg Val Val Arg Phe Trp Phe Leu
1 5 10 15
His Gln Met Val Leu Asp Leu Cys Leu Gly Gln Gly Val Pro Gln Gln
20 25 30
Asn Leu Glu Asn Pro Arg Glu Arg Lys Ser Phe Leu Leu Phe Val Arg
35 40 45

Asn Leu Ile Ile Asp Ser Ser Leu Lys Ile Leu Ser Gln Glu Pro Ser
50 55 60
Asn Leu Trp Gln Arg Ile Pro Lys Met Met Thr Thr *
65 70 75 76

<210> 1515
<211> 148
<212> PRT
<213> Homo sapiens

<400> 1515
Met Leu Gly Ser Arg Leu Met Thr Leu Thr Val Cys Ala Gly Ala Leu
1 5 10 15
Ala Arg Gly Arg Gly Thr Gly Thr Cys Glu Thr Arg Gln Glu Gly Lys
20 25 30
Gly Gln Asn His Ser Thr Leu Ala Trp Pro His Glu Glu Pro Gly Ala
35 40 45
Ser Thr Gly Arg Asp Gly Gly Lys Leu Pro Arg Gly Gln Cys Leu Leu
50 55 60
Glu Lys Gly Pro Gly Gly Ala Gly Asp Lys Val Ser Lys Ile Phe Pro
65 70 75 80
Ser Cys Ala Leu Ala Leu Leu Leu Ser Leu Ala Asn Pro Gly Pro Arg
85 90 95
Gly Pro Arg Glu Phe His Leu Cys Trp Gly Trp Leu Asp Arg Gly Val
100 105 110
Thr Gln Glu Ala Val His Val Gly Glu Lys Arg Gly Gly Leu Gly Ser
115 120 125
Gly Arg Lys Gly Gly Trp Trp Pro Gly Trp Asp Pro Gly Cys Arg Asp
130 135 140
Val Ile Thr *
145 147

<210> 1516
<211> 274
<212> PRT
<213> Homo sapiens

<400> 1516
Met Arg Gly Ser Gln Glu Val Leu Leu Met Trp Leu Leu Val Leu Ala
1 5 10 15
Val Gly Gly Thr Glu His Ala Tyr Arg Pro Gly Arg Arg Val Cys Ala
20 25 30
Val Arg Ala His Gly Asp Pro Val Ser Glu Ser Phe Val Gln Arg Val
35 40 45
Tyr Gln Pro Phe Leu Thr Thr Cys Asp Gly His Arg Ala Cys Ser Thr
50 55 60
Tyr Arg Thr Ile Tyr Arg Thr Ala Tyr Arg Arg Ser Pro Gly Leu Ala
65 70 75 80
Pro Ala Arg Pro Arg Tyr Ala Cys Cys Pro Gly Trp Lys Arg Thr Ser
85 90 95
Gly Leu Pro Gly Ala Cys Gly Ala Ala Ile Cys Gln Pro Pro Cys Arg
100 105 110
Asn Gly Gly Ser Cys Val Gln Pro Gly Arg Cys Arg Cys Pro Ala Gly

```
        115                 120                 125
Trp Arg Gly Asp Thr Cys Gln Ser Asp Val Asp Glu Cys Ser Ala Arg
    130                 135                 140
Arg Gly Gly Cys Pro Gln Arg Cys Val Asn Thr Ala Gly Ser Tyr Trp
145                 150                 155                 160
Cys Gln Cys Trp Glu Gly His Ser Leu Ser Ala Asp Gly Thr Leu Cys
                165                 170                 175
Val Pro Lys Gly Gly Pro Pro Arg Val Ala Pro Asn Pro Thr Gly Val
            180                 185                 190
Asp Ser Ala Met Lys Glu Glu Val Gln Arg Leu Gln Ser Arg Val Asp
        195                 200                 205
Leu Leu Glu Glu Lys Leu Gln Leu Val Leu Ala Pro Leu His Ser Leu
    210                 215                 220
Ala Ser Gln Ala Leu Glu His Gly Leu Pro Asp Pro Gly Ser Leu Leu
225                 230                 235                 240
Val His Ser Phe Gln Gln Leu Gly Arg Ile Asp Ser Leu Ser Glu Gln
                245                 250                 255
Ile Ser Phe Leu Glu Glu Gln Leu Gly Ser Cys Ser Cys Lys Lys Asp
            260                 265                 270
Ser  *
273


     &lt;210&gt; 1517
     &lt;211&gt; 246
     &lt;212&gt; PRT
     &lt;213&gt; Homo sapiens


     &lt;400&gt; 1517
Met Thr Leu Phe Pro Val Leu Leu Phe Leu Val Ala Gly Leu Leu Pro
  1               5                  10                  15
Ser Phe Pro Ala Asn Glu Asp Lys Asp Pro Ala Phe Thr Ala Leu Leu
            20                  25                  30
Thr Thr Gln Thr Gln Val Gln Arg Glu Ile Val Asn Lys His Asn Glu
        35                  40                  45
Leu Arg Arg Ala Val Ser Pro Pro Ala Arg Asn Met Leu Lys Met Glu
    50                  55                  60
Trp Asn Lys Glu Ala Ala Ala Asn Ala Gln Lys Trp Ala Asn Gln Cys
 65                  70                  75                  80
Asn Tyr Arg His Ser Asn Pro Lys Asp Arg Met Thr Ser Leu Lys Cys
                85                  90                  95
Gly Glu Asn Leu Tyr Met Ser Ser Ala Ser Ser Ser Trp Ser Gln Ala
            100                 105                 110
Ile Gln Ser Trp Phe Asp Glu Tyr Asn Asp Phe Asp Phe Gly Val Gly
        115                 120                 125
Pro Lys Thr Pro Asn Ala Val Val Gly His Tyr Thr Gln Val Val Trp
    130                 135                 140
Tyr Ser Ser Tyr Leu Val Gly Cys Gly Asn Ala Tyr Cys Pro Asn Gln
145                 150                 155                 160
Lys Val Leu Lys Tyr Tyr Tyr Val Cys Gln Tyr Cys Pro Ala Gly Asn
                165                 170                 175
Trp Ala Asn Arg Leu Tyr Val Pro Tyr Glu Gln Gly Ala Pro Cys Ala
            180                 185                 190
Ser Cys Pro Asp Asn Cys Asp Asp Gly Leu Cys Thr Asn Gly Cys Lys
        195                 200                 205
Tyr Glu Asp Leu Tyr Ser Asn Cys Lys Ser Leu Lys Leu Thr Leu Thr
    210                 215                 220
```

Cys Lys His Gln Leu Val Arg Asp Ser Cys Lys Ala Ser Cys Asn Cys
225 230 235 240
Ser Asn Ser Ile Tyr *
245

<210> 1518
<211> 122
<212> PRT
<213> Homo sapiens

<400> 1518
Met Arg Asn Arg Arg Thr Glu Arg Thr Cys Thr Pro Pro Leu Ala Ser
1 5 10 15
Pro Tyr Asn Leu Val Pro His Leu Gln Asn Leu Leu Ala Val Leu Leu
20 25 30
Met Ile Leu Val Leu Thr Pro Met Val Leu Asn Pro His Lys Leu Tyr
35 40 45
Gln Met Met Thr Gln Asn Ile Leu Leu Gln Lys Pro Gln Lys Asn Phe
50 55 60
Ile Trp Thr Ala Leu Lys Gly Asn Leu Ser Tyr Pro Arg Asn Leu Leu
65 70 75 80
Leu Gln Ser His Leu Ser Leu Leu Leu His Ser Leu Leu Leu Glu Leu
85 90 95
Asn Gln Arg Val Cys Leu Leu Pro Arg Ser Leu Ile Asp Pro Gly Lys
100 105 110
Arg Leu Lys Lys Lys Pro Met Glu Thr Phe
115 120 122

<210> 1519
<211> 249
<212> PRT
<213> Homo sapiens

<400> 1519
Met Gly Leu Ser Ile Phe Leu Leu Leu Cys Val Leu Gly Leu Ser Gln
1 5 10 15
Ala Ala Thr Pro Lys Ile Phe Asn Gly Thr Glu Cys Gly Arg Asn Ser
20 25 30
Gln Pro Trp Gln Val Gly Leu Phe Glu Gly Thr Ser Leu Arg Cys Gly
35 40 45
Gly Val Leu Ile Asp His Arg Trp Val Leu Thr Ala Ala His Cys Ser
50 55 60
Gly Ser Arg Tyr Trp Val Arg Leu Gly Glu His Ser Leu Ser Gln Leu
65 70 75 80
Asp Trp Thr Glu Gln Ile Arg His Ser Gly Phe Ser Val Thr His Pro
85 90 95
Gly Tyr Leu Gly Ala Ser Thr Ser His Glu His Asp Leu Arg Leu Leu
100 105 110
Arg Leu Arg Leu Pro Val Arg Val Thr Ser Ser Val Gln Pro Leu Pro
115 120 125
Leu Pro Asn Asp Cys Ala Thr Ala Gly Thr Glu Cys His Val Ser Gly
130 135 140
Trp Gly Ile Thr Asn His Pro Arg Asn Pro Phe Pro Asp Leu Leu Gln

```
145                 150                 155                 160
Cys Leu Asn Leu Ser Ile Val Ser His Ala Thr Cys His Gly Val Tyr
                165                 170                 175
Pro Gly Arg Ile Thr Ser Asn Met Val Cys Ala Gly Gly Val Pro Gly
            180                 185                 190
Gln Asp Ala Cys Gln Gly Asp Ser Gly Gly Pro Leu Val Cys Gly Gly
        195                 200                 205
Val Leu Gln Gly Leu Val Ser Trp Gly Ser Val Gly Pro Cys Gly Gln
    210                 215                 220
Asp Gly Ile Pro Gly Val Tyr Thr Tyr Ile Cys Lys Tyr Val Asp Trp
225                 230                 235                 240
Ile Arg Met Ile Met Arg Asn Asn  *
                245         248
```

```
<210> 1520
<211> 292
<212> PRT
<213> Homo sapiens


<400> 1520
Met Leu Val Leu Gln Ile Leu Leu Cys Ile Arg Glu Phe Ile Leu Glu
 1               5                  10                  15
Arg Ser Leu Ile Asn Val Lys Asn Val Ala Lys Ser Leu Ala Val Val
            20                  25                  30
Leu Ala Leu Leu Asn Ile Gly Lys Phe Ile Leu Glu Lys Ile Phe Thr
        35                  40                  45
Asn Ala Lys Tyr Val Leu Asn Leu Leu Leu Val Ser Gln Ile Leu Leu
    50                  55                  60
Cys Met Arg Glu Phe Ile Leu Glu Arg Asn Pro Ile Asn Val Lys Asn
65                  70                  75                  80
Val Ala Lys Pro Phe Leu Ile Val His Thr Leu Phe Asp Ile Ile Glu
                85                  90                  95
Phe Ile Leu Glu Lys Asn His Thr Asn Val Lys His Val Ala Asn Leu
            100                 105                 110
Leu Val Thr Pro Gln Val Leu Leu Cys Ile Gly Glu Leu Ile Leu Glu
        115                 120                 125
Arg Asn Pro Ile His Val Lys Asn Val Ala Lys Pro Leu Val Ile Val
    130                 135                 140
Gln Met Leu Phe Ser Ile Gly Glu Phe Ile Leu Ala Arg Asp Pro Thr
145                 150                 155                 160
Asn Val Lys Asn Val Ala Lys Pro Ser Thr Ile Gly His Thr Ser Leu
                165                 170                 175
His Ile Lys Glu Val Ile Leu Glu Arg Asp Pro Thr Asn Val Lys Asn
            180                 185                 190
Val Ala Lys Pro Ser Thr Leu Gly His Thr Ser Leu His Ile Gly Glu
        195                 200                 205
Asp Ile Leu Glu Arg Asp Pro Thr Asn Val Met Asn Val Val Lys Pro
    210                 215                 220
Ser Ala Ile Gly His Thr Ser Leu His Ile Gly Glu Val Ile Val Glu
225                 230                 235                 240
Arg Asp Pro Thr Asn Val Lys Asn Val Ala Lys Pro Leu Thr Leu Gly
                245                 250                 255
His Thr Ser Leu His Ile Arg Glu Val Ile Leu Glu Lys Asn Phe Lys
            260                 265                 270
Asn Val Lys His Gly Ala Asp Phe Leu Leu Val Thr His Val Leu Leu
        275                 280                 285
```

```
Cys Ile Arg  *
    290 291


     <210> 1521
     <211> 129
     <212> PRT
     <213> Homo sapiens


     <400> 1521
Met Gly Ser Thr Ala Ile Leu Ala Leu Leu Leu Ala Val Leu Gln Gly
  1               5                  10                  15
Val Cys Ala Glu Val Gln Leu Val Gln Ser Gly Ala Glu Val Lys Lys
             20                  25                  30
Pro Gly Glu Ser Leu Lys Ile Ser Cys Lys Gly Ser Gly Tyr Ser Phe
         35                  40                  45
Thr Ser Tyr Trp Ile Gly Trp Val Arg Gln Met Pro Gly Lys Gly Leu
     50                  55                  60
Glu Trp Met Gly Ile Ile Tyr Pro Gly Asp Ser Asp Thr Arg Tyr Ser
 65                  70                  75                  80
Pro Ser Phe Gln Gly Gln Val Thr Ile Ser Ala Asp Lys Ser Ile Ser
                 85                  90                  95
Thr Ala Tyr Leu Gln Trp Ser Ser Leu Lys Ala Ser Asp Thr Ala Met
            100                 105                 110
Tyr Tyr Cys Ala Arg His Thr Val Arg Glu Thr Ser Pro Glu Pro Val
        115                 120                 125             128
 *


     <210> 1522
     <211> 66
     <212> PRT
     <213> Homo sapiens


     <400> 1522
Met Val Val Val Leu Pro Cys Phe Ala Val Leu Lys Leu Leu Phe Gly
  1               5                  10                  15
Gln Ser Lys Leu Gly Pro Met Gln Pro Ser Gln Ser Gly Leu Asp Pro
             20                  25                  30
Val Gly Ala Gly Met Ser Ala Ser Ile Ala Asp Gly Ser Arg Ala Thr
         35                  40                  45
Ala Asp Lys Ala Val Leu Leu Asp Pro Thr Ser Leu Leu Leu Glu Tyr
     50                  55                  60
Thr  *
 65


     <210> 1523
     <211> 131
     <212> PRT
     <213> Homo sapiens
```

<400> 1523
```
Met Ile Leu Leu Ala Phe Leu Val Cys Trp Gly Pro Leu Phe Gly Leu
 1               5                  10                  15
Leu Leu Ala Asp Val Phe Gly Ser Asn Leu Trp Ala Gln Glu Tyr Leu
            20                  25                  30
Arg Gly Met Asp Trp Ile Leu Ala Leu Ala Val Leu Asn Ser Ala Val
        35                  40                  45
Asn Pro Ile Ile Tyr Ser Phe Arg Ser Arg Glu Val Cys Arg Ala Val
    50                  55                  60
Leu Ser Phe Leu Cys Cys Gly Cys Leu Arg Leu Gly Met Arg Gly Pro
65                  70                  75                  80
Gly Asp Cys Leu Ala Arg Ala Val Glu Ala His Ser Gly Ala Ser Thr
                85                  90                  95
Thr Asp Ser Ser Leu Arg Pro Arg Asp Ser Phe Arg Gly Ser Arg Ser
            100                 105                 110
Leu Ser Phe Arg Met Arg Glu Pro Leu Ser Ser Ile Ser Ser Val Arg
        115                 120                 125
Ser Ile  *
    130
```

<210> 1524
<211> 52
<212> PRT
<213> Homo sapiens

<400> 1524
```
Met Lys Phe Phe Val Phe Ala Leu Ile Leu Ala Leu Met Leu Ser Met
 1               5                  10                  15
Thr Gly Ala Asp Ser His Ala Lys Arg His His Gly Tyr Lys Arg Lys
            20                  25                  30
Phe His Glu Lys His His Ser His Arg Gly Tyr Arg Ser Asn Tyr Leu
        35                  40                  45
Tyr Asp Asn  *
    50  51
```

<210> 1525
<211> 246
<212> PRT
<213> Homo sapiens

<400> 1525
```
Met Thr Leu Phe Pro Val Leu Leu Phe Leu Val Ala Gly Leu Leu Pro
 1               5                  10                  15
Ser Phe Pro Ala Asn Glu Asp Lys Asp Pro Ala Phe Thr Ala Leu Leu
            20                  25                  30
Thr Thr Gln Thr Gln Val Gln Arg Glu Ile Val Asn Lys His Asn Glu
        35                  40                  45
Leu Arg Arg Ala Val Ser Pro Pro Ala Arg Asn Met Leu Lys Met Glu
    50                  55                  60
Trp Asn Lys Glu Ala Ala Ala Asn Ala Gln Lys Trp Ala Asn Gln Cys
65                  70                  75                  80
Asn Tyr Arg His Ser Asn Pro Lys Asp Arg Met Thr Ser Leu Lys Cys
                85                  90                  95
```

Gly Glu Asn Leu Tyr Met Ser Ser Ala Ser Ser Ser Trp Ser Gln Ala
100 105 110
Ile Gln Ser Trp Phe Asp Glu Tyr Asn Asp Phe Asp Phe Gly Val Gly
115 120 125
Pro Lys Thr Pro Asn Ala Val Val Gly His Tyr Thr Gln Val Val Trp
130 135 140
Tyr Ser Ser Tyr Leu Val Gly Cys Gly Asn Ala Tyr Cys Pro Asn Gln
145 150 155 160
Lys Val Leu Lys Tyr Tyr Tyr Val Cys Gln Tyr Cys Pro Ala Gly Asn
165 170 175
Trp Ala Asn Arg Leu Tyr Val Pro Tyr Glu Gln Gly Ala Pro Cys Ala
180 185 190
Ser Cys Pro Asp Asn Cys Asp Asp Gly Leu Cys Thr Asn Gly Cys Lys
195 200 205
Tyr Glu Asp Leu Tyr Ser Asn Cys Lys Ser Leu Lys Leu Thr Leu Thr
210 215 220
Cys Lys His Gln Leu Val Arg Asp Ser Cys Lys Ala Ser Cys Asn Cys
225 230 235 240
Ser Asn Ser Ile Tyr *
245

<210> 1526
<211> 47
<212> PRT
<213> Homo sapiens

<400> 1526
Met Val Leu Gly Ala Arg Ala Val Ile Ser Phe Cys Ile Leu Ser Ala
1 5 10 15
Met Pro Gly Tyr Met Val Val Pro Pro Glu Arg Thr Leu Leu Ala Tyr
20 25 30
Lys Ser Leu Arg Met Ser Met Ser His Phe Met Met Glu Leu *
35 40 45 46

<210> 1527
<211> 118
<212> PRT
<213> Homo sapiens

<400> 1527
Met Ser Ala Arg Gly Trp Pro Cys Glu Ala Phe Val Leu Ala Gln Val
1 5 10 15
Cys Trp Cys Trp Leu Cys Val Arg Gly Arg Leu Cys Glu Ala Leu Thr
20 25 30
Leu Ala Gln Val Arg Arg His Gln Val Cys Val Pro Gly Gln Pro Cys
35 40 45
Glu Ala Leu Thr Leu Thr Gln Val Arg Arg His Gln Leu Cys Val Trp
50 55 60
Gly Arg Pro Cys Glu Ala Leu Thr Leu Ala Gln Val Cys Trp Leu Trp
65 70 75 80
Leu Cys Val Gln Gly Trp Pro His Glu Ala Leu Thr Leu Ala Gln Val
85 90 95
Arg Gln His Gln Val Cys Val Arg Gly Arg Pro Cys Glu Ala Leu Ser

100 105 110
Leu Ala Gln Val Arg *
115 117

<210> 1528
<211> 92
<212> PRT
<213> Homo sapiens

<400> 1528
Met Lys Val Ser Ala Ala Ala Leu Ala Val Ile Leu Ile Ala Thr Ala
1 5 10 15
Leu Cys Ala Pro Ala Ser Ala Ser Pro Tyr Ser Ser Asp Thr Thr Pro
20 25 30
Cys Cys Phe Ala Tyr Ile Ala Arg Pro Leu Pro Arg Ala His Ile Lys
35 40 45
Glu Tyr Phe Tyr Thr Ser Gly Lys Cys Ser Asn Pro Ala Val Val Phe
50 55 60
Val Thr Arg Lys Asn Arg Gln Val Cys Ala Asn Pro Glu Lys Lys Trp
65 70 75 80
Val Arg Glu Tyr Ile Asn Ser Leu Glu Met Ser *
85 90 91

<210> 1529
<211> 71
<212> PRT
<213> Homo sapiens

<400> 1529
Met Tyr Cys Trp Trp Cys Trp Leu Cys Thr Ala Met Val Cys Ser Gly
1 5 10 15
Val Leu Cys Arg Pro Leu Trp Glu Pro Leu Ser Pro Arg Leu Ser Val
20 25 30
Phe Trp Ala Gly Arg Tyr Leu Gly Phe Trp Cys Met Gly Cys Cys Arg
35 40 45
Met Ala Met Tyr Cys Val Ser Ser Cys Ser Arg Phe Ser Gly Glu Ser
50 55 60
Gly Phe Arg Arg Ile Pro *
65 70

<210> 1530
<211> 85
<212> PRT
<213> Homo sapiens

<400> 1530
Met Val Leu Arg Val Cys Phe Leu Ile Phe Val Leu Tyr His Asn Leu
1 5 10 15
Gly Lys Tyr Ile Phe Ile Ile Tyr Val Tyr Arg Cys Lys Asp Arg Phe
20 25 30

Thr Lys Gly Cys Ile Thr Val Val Gln Gln Ser Gly Ile Leu Thr Glu
35 40 45
Leu Lys Gly Gln Gly Ser Phe Leu Tyr Val Leu Leu Cys Leu Asp Ile
50 55 60
Thr Leu Leu Val Arg Ser Val Phe Lys Asn Asp Asn Ser Arg Phe Asp
65 70 75 80
Phe Gln Ala Asn *
84

<210> 1531
<211> 60
<212> PRT
<213> Homo sapiens

<400> 1531
Met Leu Pro Gln Val Phe Leu Gly Phe Thr Lys Val Arg Leu Leu Arg
1 5 10 15
Leu Arg Asn Pro Trp Gly Cys Val Glu Trp Thr Gly Ala Trp Ser Asp
20 25 30
Arg Trp Asp Gly Ser Gly Val Gly Val Gly Leu Asp Pro Thr Cys Pro
35 40 45
Pro Leu Thr Pro Gln Ser Leu Gln Leu Pro Thr Leu
50 55 60

<210> 1532
<211> 53
<212> PRT
<213> Homo sapiens

<400> 1532
Met Leu Gly Leu His Gln Leu Cys Ser Leu Leu Val Gln Leu Asp Phe
1 5 10 15
Tyr Leu Gln Tyr Leu Tyr Gly Gln Phe Gln Gln Phe Ser Met Cys Leu
20 25 30
Asp Leu Asn His Val His Phe Leu Met Phe Pro Ser Leu Val Cys Ala
35 40 45
Met Phe Arg Phe *
50 52

<210> 1533
<211> 741
<212> PRT
<213> Homo sapiens

<400> 1533
Met Ala Glu Ser Arg Gly Arg Leu Tyr Leu Trp Met Cys Leu Ala Ala
1 5 10 15
Ala Leu Ala Ser Phe Leu Met Gly Phe Met Val Gly Trp Phe Ile Lys
20 25 30
Pro Leu Lys Glu Thr Thr Thr Ser Val Arg Tyr His Gln Ser Ile Arg

35 40 45
Trp Lys Leu Val Ser Glu Met Lys Ala Glu Asn Ile Lys Ser Phe Leu
50 55 60
Arg Ser Phe Thr Lys Leu Pro His Leu Ala Gly Thr Glu Gln Asn Phe
65 70 75 80
Leu Leu Ala Lys Lys Ile Gln Thr Gln Trp Lys Lys Phe Gly Leu Asp
85 90 95
Ser Ala Lys Leu Val His Tyr Asp Val Leu Leu Ser Tyr Pro Asn Glu
100 105 110
Thr Asn Ala Asn Tyr Ile Ser Ile Val Asp Glu His Glu Thr Glu Ile
115 120 125
Phe Lys Thr Ser Tyr Leu Glu Pro Pro Pro Asp Gly Tyr Glu Asn Val
130 135 140
Thr Asn Ile Val Pro Pro Tyr Asn Ala Phe Ser Ala Gln Gly Met Pro
145 150 155 160
Glu Gly Asp Leu Val Tyr Val Asn Tyr Ala Arg Thr Glu Asp Phe Phe
165 170 175
Lys Leu Glu Arg Glu Met Gly Ile Asn Cys Thr Gly Lys Ile Val Ile
180 185 190
Ala Arg Tyr Gly Lys Ile Phe Arg Gly Asn Lys Val Lys Asn Ala Met
195 200 205
Leu Ala Gly Ala Ile Gly Ile Ile Leu Tyr Ser Asp Pro Ala Asp Tyr
210 215 220
Phe Ala Pro Glu Val Gln Pro Tyr Pro Lys Gly Trp Asn Leu Pro Gly
225 230 235 240
Thr Ala Ala Gln Arg Gly Asn Val Leu Asn Leu Asn Gly Ala Gly Asp
245 250 255
Pro Leu Thr Pro Gly Tyr Pro Ala Lys Glu Tyr Thr Phe Arg Leu Asp
260 265 270
Val Glu Glu Gly Val Gly Ile Pro Arg Ile Pro Val His Pro Ile Gly
275 280 285
Tyr Asn Asp Ala Glu Ile Leu Leu Arg Tyr Leu Gly Gly Ile Ala Pro
290 295 300
Pro Asp Lys Ser Trp Lys Gly Ala Leu Asn Val Ser Tyr Ser Ile Gly
305 310 315 320
Pro Gly Phe Thr Gly Ser Asp Ser Phe Arg Lys Val Arg Met His Val
325 330 335
Tyr Asn Ile Asn Lys Ile Thr Arg Ile Tyr Asn Val Val Gly Thr Ile
340 345 350
Arg Gly Ser Val Glu Pro Asp Arg Tyr Val Ile Leu Gly Gly His Arg
355 360 365
Asp Ser Trp Val Phe Gly Ala Ile Asp Pro Thr Ser Gly Val Ala Val
370 375 380
Leu Gln Glu Ile Ala Arg Ser Phe Gly Lys Leu Met Ser Lys Gly Trp
385 390 395 400
Arg Pro Arg Arg Thr Ile Ile Phe Ala Ser Trp Asp Ala Glu Glu Phe
405 410 415
Gly Leu Leu Gly Ser Thr Glu Trp Ala Glu Glu Asn Val Lys Ile Leu
420 425 430
Gln Glu Arg Ser Ile Ala Tyr Ile Asn Ser Asp Ser Ser Ile Glu Gly
435 440 445
Asn Tyr Thr Leu Arg Val Asp Cys Thr Pro Leu Leu Tyr Gln Leu Val
450 455 460
Tyr Lys Leu Thr Lys Glu Ile Pro Ser Pro Asp Asp Gly Phe Glu Ser
465 470 475 480
Lys Phe Leu Tyr Glu Ser Trp Val Glu Lys Asp Pro Ser Pro Glu Asn
485 490 495
Lys Asn Leu Pro Arg Ile Asn Lys Leu Gly Ser Gly Ser Asp Phe Glu
500 505 510

Ala Tyr Phe Gln Arg Leu Gly Ile Ala Ser Gly Arg Ala Arg Tyr Thr
515 520 525
Lys Asn Lys Lys Thr Asp Lys Tyr Ser Ser Tyr Pro Val Tyr His Thr
530 535 540
Ile Tyr Glu Thr Phe Glu Leu Val Glu Lys Phe Tyr Asp Pro Thr Phe
545 550 555 560
Lys Lys Gln Leu Ser Val Ala Gln Leu Arg Gly Ala Leu Val Tyr Glu
565 570 575
Leu Val Asp Ser Lys Ile Ile Pro Phe Asn Ile Gln Asp Tyr Ala Glu
580 585 590
Ala Leu Lys Asn Tyr Ala Ala Ser Ile Tyr Asn Leu Ser Lys Lys His
595 600 605
Asp Gln Gln Leu Thr Asp His Gly Val Ser Phe Asp Ser Leu Phe Ser
610 615 620
Ala Val Lys Asn Phe Ser Glu Ala Ala Ser Asp Phe His Lys Arg Leu
625 630 635 640
Ile Gln Val Asp Leu Asn Asn Pro Ile Ala Val Arg Met Met Asn Asp
645 650 655
Gln Leu Met Leu Leu Glu Arg Ala Phe Ile Asp Pro Leu Gly Leu Pro
660 665 670
Gly Lys Leu Phe Tyr Arg His Ile Ile Phe Ala Pro Ser Ser His Asn
675 680 685
Lys Tyr Ala Gly Glu Ser Phe Pro Gly Ile Tyr Asp Ala Ile Phe Asp
690 695 700
Ile Glu Asn Lys Ala Asn Ser Arg Leu Ala Trp Lys Glu Val Lys Lys
705 710 715 720
His Ile Ser Ile Ala Ala Phe Thr Ile Gln Ala Ala Ala Gly Thr Leu
725 730 735
Lys Glu Val Leu *
740

<210> 1534
<211> 50
<212> PRT
<213> Homo sapiens

<400> 1534
Met Leu Ile Leu Leu His Ile Leu Lys Asn Ile Lys Leu Tyr Leu Val
1 5 10 15
Asn Met Leu Lys Thr Lys Leu Cys Phe Tyr Lys Asp Arg Gly Ser Pro
20 25 30
Glu Glu Gly Ile Asp Lys Glu Glu Met Lys Leu Gly Gly Arg Lys Trp
35 40 45
Thr *
49

<210> 1535
<211> 973
<212> PRT
<213> Homo sapiens

<400> 1535
Met Val Lys Ser Lys Trp Gly Leu Ala Leu Ala Ala Val Val Thr Val

```
 1               5                   10                  15
Leu Ser Ser Leu Leu Met Ser Val Gly Leu Cys Thr Leu Phe Gly Leu
            20                  25                  30
Thr Pro Thr Leu Asn Gly Gly Glu Ile Phe Pro Tyr Leu Val Val Val
        35                  40                  45
Ile Gly Leu Glu Asn Val Leu Val Leu Thr Lys Ser Val Val Ser Thr
    50                  55                  60
Pro Val Asp Leu Glu Val Lys Leu Arg Ile Ala Gln Gly Leu Ser Ser
65                  70                  75                  80
Glu Ser Trp Ser Ile Met Lys Asn Met Ala Thr Glu Leu Gly Ile Ile
                85                  90                  95
Leu Ile Gly Tyr Phe Thr Leu Val Pro Ala Ile Gln Glu Phe Cys Leu
            100                 105                 110
Phe Ala Val Val Gly Leu Val Ser Asp Phe Phe Leu Gln Met Leu Phe
        115                 120                 125
Phe Thr Thr Val Leu Ser Ile Asp Ile Arg Arg Met Glu Leu Ala Asp
    130                 135                 140
Leu Asn Lys Arg Leu Pro Pro Glu Ala Cys Leu Pro Ser Ala Lys Pro
145                 150                 155                 160
Val Gly Gln Pro Thr Arg Tyr Glu Arg Gln Leu Ala Val Arg Pro Ser
                165                 170                 175
Thr Pro His Thr Ile Thr Leu Gln Pro Ser Ser Phe Arg Asn Leu Arg
            180                 185                 190
Leu Pro Lys Arg Leu Arg Val Val Tyr Phe Leu Ala Arg Thr Arg Leu
        195                 200                 205
Ala Gln Arg Leu Ile Met Ala Gly Thr Val Val Trp Ile Gly Ile Leu
    210                 215                 220
Val Tyr Thr Asp Pro Ala Gly Leu Arg Asn Tyr Leu Ala Ala Gln Val
225                 230                 235                 240
Thr Glu Gln Ser Pro Leu Gly Glu Gly Ala Leu Ala Pro Met Pro Val
                245                 250                 255
Pro Ser Gly Met Leu Pro Pro Ser His Pro Asp Pro Ala Phe Ser Ile
            260                 265                 270
Phe Pro Pro Asp Ala Pro Lys Leu Pro Glu Asn Gln Thr Ser Pro Gly
        275                 280                 285
Glu Ser Pro Glu Arg Gly Gly Pro Ala Glu Val Val His Asp Ser Pro
    290                 295                 300
Val Pro Glu Val Thr Trp Gly Pro Glu Asp Glu Glu Leu Trp Arg Lys
305                 310                 315                 320
Leu Ser Phe Arg His Trp Pro Thr Leu Phe Ser Tyr Tyr Asn Ile Thr
                325                 330                 335
Leu Ala Lys Arg Tyr Ile Ser Leu Leu Pro Val Ile Pro Val Thr Leu
            340                 345                 350
Arg Leu Asn Pro Arg Glu Ala Leu Glu Gly Arg His Pro Gln Asp Gly
        355                 360                 365
Arg Ser Ala Trp Pro Pro Pro Gly Pro Ile Pro Ala Gly His Trp Glu
    370                 375                 380
Ala Gly Pro Lys Gly Pro Gly Gly Val Gln Ala His Gly Asp Val Thr
385                 390                 395                 400
Leu Tyr Lys Val Ala Ala Leu Gly Leu Ala Thr Gly Ile Val Leu Val
                405                 410                 415
Leu Leu Leu Leu Cys Leu Tyr Arg Val Leu Cys Pro Arg Asn Tyr Gly
            420                 425                 430
Gln Leu Gly Gly Gly Pro Gly Arg Arg Arg Arg Gly Glu Leu Pro Cys
        435                 440                 445
Asp Asp Tyr Gly Tyr Ala Pro Pro Glu Thr Glu Ile Val Pro Leu Val
    450                 455                 460
Leu Arg Gly His Leu Met Asp Ile Glu Cys Leu Ala Ser Asp Gly Met
465                 470                 475                 480
```

Leu Leu Val Ser Cys Cys Leu Ala Gly His Val Cys Val Trp Asp Ala
485 490 495
Gln Thr Gly Asp Cys Leu Thr Arg Ile Pro Arg Pro Gly Arg Gln Arg
500 505 510
Arg Asp Ser Gly Val Gly Ser Gly Leu Glu Ala Gln Glu Ser Trp Glu
515 520 525
Arg Leu Ser Asp Gly Gly Lys Ala Gly Pro Glu Glu Pro Gly Asp Ser
530 535 540
Pro Pro Leu Arg His Arg Pro Arg Gly Pro Pro Pro Pro Ser Leu Phe
545 550 555 560
Gly Asp Gln Pro Asp Leu Thr Cys Leu Ile Asp Thr Asn Phe Ser Ala
565 570 575
Gln Pro Arg Ser Ser Gln Pro Thr Gln Pro Glu Pro Arg His Arg Ala
580 585 590
Val Cys Gly Arg Ser Arg Asp Ser Pro Gly Tyr Asp Phe Ser Cys Leu
595 600 605
Val Gln Arg Val Tyr Gln Glu Glu Gly Leu Ala Ala Val Cys Thr Pro
610 615 620
Ala Leu Arg Pro Pro Ser Pro Gly Pro Val Leu Ser Gln Ala Pro Glu
625 630 635 640
Asp Glu Gly Gly Ser Pro Glu Lys Gly Ser Pro Ser Leu Ala Trp Ala
645 650 655
Pro Ser Ala Glu Gly Ser Ile Trp Ser Leu Glu Leu Gln Gly Asn Leu
660 665 670
Ile Val Val Gly Arg Ser Ser Gly Arg Leu Glu Val Trp Asp Ala Ile
675 680 685
Glu Gly Val Leu Cys Cys Ser Ser Glu Glu Val Ser Ser Gly Ile Thr
690 695 700
Ala Leu Val Phe Leu Asp Lys Arg Ile Val Ala Ala Arg Leu Asn Gly
705 710 715 720
Ser Leu Asp Phe Phe Ser Leu Glu Thr His Thr Ala Leu Ser Pro Leu
725 730 735
Gln Phe Arg Gly Thr Pro Gly Arg Gly Ser Ser Pro Ala Ser Pro Val
740 745 750
Tyr Ser Ser Ser Asp Thr Val Ala Cys His Leu Thr His Thr Val Pro
755 760 765
Cys Ala His Gln Lys Pro Ile Thr Ala Leu Lys Ala Ala Ala Gly Arg
770 775 780
Leu Val Thr Gly Ser Gln Asp His Thr Leu Arg Val Phe Arg Leu Glu
785 790 795 800
Asp Ser Cys Cys Leu Phe Thr Leu Gln Gly His Ser Gly Ala Ile Thr
805 810 815
Thr Val Tyr Ile Asp Gln Thr Met Val Leu Ala Ser Gly Gly Gln Asp
820 825 830
Gly Ala Ile Cys Leu Trp Asp Val Leu Thr Gly Ser Arg Val Ser His
835 840 845
Val Phe Ala His Arg Gly Asp Val Thr Ser Leu Thr Cys Thr Thr Ser
850 855 860
Cys Val Ile Ser Ser Gly Leu Asp Asp Leu Ile Ser Ile Trp Asp Arg
865 870 875 880
Ser Thr Gly Ile Lys Phe Tyr Ser Ile Gln Gln Asp Leu Gly Cys Gly
885 890 895
Ala Ser Leu Gly Val Ile Ser Asp Asn Leu Leu Val Thr Gly Gly Gln
900 905 910
Gly Cys Val Ser Phe Trp Asp Leu Asn Tyr Gly Asp Leu Leu Gln Thr
915 920 925
Val Tyr Leu Gly Lys Asn Ser Glu Ala Gln Pro Ala Arg Gln Ile Leu
930 935 940
Val Leu Asp Asn Ala Ala Ile Val Cys Asn Phe Gly Ser Glu Leu Ser

```
945                 950                 955                 960
Leu Val Tyr Val Pro Ser Val Leu Glu Lys Leu Asp  *
                965                 970     972
```

<210> 1536
<211> 75
<212> PRT
<213> Homo sapiens

<400> 1536

```
Met Cys Leu Leu Lys Ala Ala Pro Phe Phe Phe Phe Tyr Val Pro Gln
  1               5                  10                  15
Val Gly Lys Gly Asn Pro Arg Pro Pro Arg Gly Cys Ser Ala Phe His
             20                  25                  30
Pro Pro Thr His Leu Arg Pro Gly Ser Cys Ser Val Ala Gln Ala Gly
         35                  40                  45
Val Gln Trp Arg Ser Leu Gly Ser Ile Ala Ala Ser Val Ser Trp Val
     50                  55                  60
Gln Ala Ile Leu Leu Pro Gln Pro Leu Glu  *
 65                  70              74
```

<210> 1537
<211> 96
<212> PRT
<213> Homo sapiens

<400> 1537

```
Met Asp Leu Gly Arg Val Phe Ile Thr Leu Ile Leu Asn Leu Leu Arg
  1               5                  10                  15
Glu Thr Ile Phe Lys Arg Asp Gln Ser Pro Glu Pro Lys Val Pro Glu
             20                  25                  30
Gln Ser Val Lys Glu Asp Arg Lys Leu Cys Glu Arg Pro Leu Ala Ser
         35                  40                  45
Ser Pro Pro Arg Leu Tyr Glu Asp Asp Glu Thr Pro Gly Ala Leu Ser
     50                  55                  60
Gly Leu Thr Asn Met Ala Val Ile Gln Ile Asp Gly His Met Ser Gly
 65                  70                  75                  80
Gln Met Val Lys His Leu Met Asn Ser Met Met Lys Leu Cys Val Met
                 85                  90                  95  96
```

<210> 1538
<211> 318
<212> PRT
<213> Homo sapiens

<400> 1538

```
Met Val Met Arg Pro Leu Trp Ser Leu Leu Leu Trp Glu Ala Leu Leu
  1               5                  10                  15
```

Pro Ile Thr Val Thr Gly Ala Gln Val Leu Ser Lys Val Gly Gly Ser
20 25 30
Val Leu Leu Val Ala Ala Arg Pro Pro Gly Phe Gln Val Arg Glu Ala
35 40 45
Ile Trp Arg Ser Leu Trp Pro Ser Glu Glu Leu Leu Ala Thr Phe Phe
50 55 60
Arg Gly Ser Leu Glu Thr Leu Tyr His Ser Arg Phe Leu Gly Arg Ala
65 70 75 80
Gln Leu His Ser Asn Leu Ser Leu Glu Leu Gly Pro Leu Glu Ser Gly
85 90 95
Asp Ser Gly Asn Phe Ser Val Leu Met Val Asp Thr Arg Gly Gln Pro
100 105 110
Trp Thr Gln Thr Leu Gln Leu Lys Val Tyr Asp Ala Val Pro Arg Pro
115 120 125
Val Val Gln Val Phe Ile Ala Val Glu Arg Asp Ala Gln Pro Ser Lys
130 135 140
Thr Cys Gln Val Phe Leu Ser Cys Trp Ala Pro Asn Ile Ser Glu Ile
145 150 155 160
Thr Tyr Ser Trp Arg Arg Glu Thr Thr Met Asp Phe Gly Met Glu Pro
165 170 175
His Ser Leu Phe Thr Asp Gly Gln Val Leu Ser Ile Ser Leu Gly Pro
180 185 190
Gly Asp Arg Asp Val Ala Tyr Ser Cys Ile Val Ser Asn Pro Val Ser
195 200 205
Trp Asp Leu Ala Thr Val Thr Pro Trp Asp Ser Cys His His Glu Ala
210 215 220
Ala Pro Gly Lys Ala Ser Tyr Lys Asp Val Leu Leu Val Val Val Pro
225 230 235 240
Val Ser Leu Leu Leu Met Leu Val Thr Leu Phe Ser Ala Trp His Trp
245 250 255
Cys Pro Cys Ser Gly Pro His Leu Arg Ser Lys Gln Leu Trp Met Arg
260 265 270
Trp Asp Leu Gln Leu Ser Leu His Lys Val Thr Leu Ser Asn Leu Ile
275 280 285
Ser Thr Val Val Cys Ser Val Val His Gln Gly Leu Val Glu Gln Ile
290 295 300
His Thr Ala Leu Ile Lys Phe Pro Ser Leu Met Lys Lys Lys
305 310 315 318

<210> 1539
<211> 157
<212> PRT
<213> Homo sapiens

<400> 1539
Met Ile Leu Gln Val Ser Gly Gly Pro Trp Thr Val Ala Leu Thr Ala
1 5 10 15
Leu Leu Met Val Leu Leu Ile Ser Val Val Gln Ser Arg Ala Thr Pro
20 25 30
Glu Asn Ser Val Tyr Gln Glu Arg Gln Glu Cys Tyr Ala Phe Asn Gly
35 40 45
Thr Gln Arg Val Val Asp Gly Leu Ile Tyr Asn Arg Glu Glu Tyr Val
50 55 60
His Phe Asp Ser Ala Val Gly Glu Phe Leu Ala Val Met Glu Leu Gly
65 70 75 80
Arg Pro Ile Gly Glu Tyr Phe Asn Ser Gln Lys Asp Phe Met Glu Arg

```
                85                      90                      95
Lys Arg Ala Glu Val Asp Lys Val Cys Arg His Lys Tyr Glu Leu Met
            100                     105                     110
Glu Pro Leu Ile Arg Gln Arg Arg Gly Asp Val Thr Ile Thr Ala Val
        115                     120                     125
Arg Gly Cys Trp Thr Thr Ile Leu Ser Gly Tyr Phe Leu Leu Lys Arg
    130                     135                     140
Gly Val Val Ser Gly Gly Cys Ser Trp Gly Ser Ser  *
145                 150                     155 156
```

<210> 1540
<211> 135
<212> PRT
<213> Homo sapiens

<400> 1540

```
Met Gly Ser Ser Phe Ile Leu Ala Leu Leu Leu Ala Val Leu Gln Gly
 1               5                  10                  15
Leu Ser Ala Gly Val Leu Leu Glu Gln Ser Arg Ala Glu Val Lys Lys
            20                  25                  30
Pro Gly Glu Ser Leu Lys Ile Ser Cys Lys Ala Ser Gly Tyr Arg Phe
        35                  40                  45
Thr Ser Ala Trp Ile Ala Trp Val Arg Gln Met Pro Gly Lys Gly Leu
    50                  55                  60
Glu Trp Met Gly Thr Ile Tyr Pro Ala Asp Ser Glu Val Arg Tyr Ser
65                  70                  75                  80
Pro Ser Leu Gln Gly Gln Val Thr Leu Ser Val Asp Glu Ser Ile Ser
                85                  90                  95
Thr Ala Tyr Leu Gln Trp Asn Ser Leu Arg Ala Ser Asp Thr Ala Thr
            100                 105                 110
Tyr Tyr Cys Ala Arg Gln Ile Ile Gly Ala Leu Pro Thr Asp Pro Phe
        115                 120                 125
Asp Leu Leu Gly Gln Gly Thr
    130                 135
```

<210> 1541
<211> 72
<212> PRT
<213> Homo sapiens

<400> 1541

```
Met Cys Val Thr Cys Val Val Cys Met Trp Cys Met Cys Gly Val Cys
 1               5                  10                  15
Ala Met Tyr Val Ala Cys Val Met His Val Val Cys Glu Val Tyr Val
            20                  25                  30
Trp Tyr Val Cys Asp Val Cys Ala Phe Gly His Thr Gly Val Val Ile
        35                  40                  45
Ala Leu Thr Trp Thr Pro Pro Gln Arg Val Ile Arg Lys Gly Gln Val
    50                  55                  60
Leu Arg Leu Ala Cys Ser Gln  *
65                  70  71
```

<210> 1542
<211> 369
<212> PRT
<213> Homo sapiens

<400> 1542
```
Met Ala Pro Arg Thr Leu Val Leu Leu Leu Ser Gly Ala Leu Ala Leu
 1               5                  10                  15
Thr Gln Thr Trp Ala Gly Ser His Ser Met Arg Tyr Phe Phe Thr Ser
            20                  25                  30
Val Ser Arg Pro Gly Arg Gly Glu Pro Arg Phe Ile Ala Val Gly Tyr
        35                  40                  45
Val Asp Asp Thr Gln Phe Val Arg Phe Asp Ser Asp Ala Ala Ser Gln
    50                  55                  60
Arg Met Glu Pro Arg Ala Pro Trp Ile Glu Gln Glu Gly Pro Glu Tyr
65                  70                  75                  80
Trp Asp Gly Glu Thr Arg Lys Val Lys Ala His Ser Gln Thr His Arg
                85                  90                  95
Val Asp Leu Gly Thr Leu Arg Gly Tyr Tyr Asn Gln Ser Glu Ala Gly
            100                 105                 110
Ser His Thr Val Gln Arg Met Tyr Gly Cys Asp Val Gly Ser Asp Trp
        115                 120                 125
Arg Phe Leu Arg Gly Tyr His Gln Tyr Ala Tyr Asp Gly Lys Asp Tyr
    130                 135                 140
Ile Ala Leu Lys Glu Asp Leu Arg Ser Trp Thr Ala Ala Asp Met Ala
145                 150                 155                 160
Ala Gln Thr Thr Lys His Lys Trp Glu Ala Ala His Val Ala Glu Gln
                165                 170                 175
Leu Arg Ala Tyr Leu Glu Gly Thr Cys Val Glu Trp Leu Arg Arg Tyr
            180                 185                 190
Leu Glu Asn Gly Lys Glu Thr Leu Gln Arg Thr Asp Ala Pro Lys Thr
        195                 200                 205
His Met Thr His His Pro Ile Ser Asp His Glu Ala Thr Leu Arg Cys
    210                 215                 220
Trp Ala Leu Ser Phe Tyr Pro Ala Glu Ile Thr Leu Thr Trp Gln Arg
225                 230                 235                 240
Asp Gly Glu Asp Gln Thr Gln Asp Thr Glu Leu Val Glu Thr Arg Pro
                245                 250                 255
Ala Gly Asp Gly Thr Phe Gln Lys Trp Ala Ala Val Val Val Pro Ser
            260                 265                 270
Gly Gln Glu Gln Arg Tyr Thr Cys His Val Gln His Glu Gly Leu Pro
        275                 280                 285
Lys Pro Leu Thr Leu Arg Trp Glu Pro Ser Ser Gln Pro Thr Ile Pro
    290                 295                 300
Ile Val Gly Ile Ile Ala Gly Leu Val Leu Phe Gly Ala Val Ile Thr
305                 310                 315                 320
Gly Ala Val Val Ala Ala Val Met Trp Arg Arg Lys Ser Ser Asp Arg
                325                 330                 335
Lys Gly Val Lys Asp Arg Lys Gly Gly Ser Tyr Ser Gln Ala Ala Ser
            340                 345                 350
Ser Asp Ser Ala Gln Gly Ser Asp Val Ser Leu Thr Ala Cys Lys Val
        355                 360                 365                 368
 *
```

<210> 1543
<211> 49
<212> PRT
<213> Homo sapiens

<400> 1543
Met Arg Ser Leu Trp Lys Ala Asn Arg Ala Asp Leu Leu Ile Trp Leu
1 5 10 15
Val Thr Phe Thr Ala Thr Ile Leu Leu Asn Leu Asp Leu Gly Leu Glu
20 25 30
Asp Ala Val Ile Phe Ser Leu Leu Leu Glu Glu Val Arg Thr Gln Met
35 40 45 48
*

<210> 1544
<211> 121
<212> PRT
<213> Homo sapiens

<400> 1544
Met Lys Ile Phe Lys Cys Tyr Phe Lys His Thr Leu Gln Gln Lys Val
1 5 10 15
Phe Ile Leu Phe Leu Thr Leu Trp Leu Leu Ser Leu Leu Lys Leu Leu
20 25 30
Asn Val Arg Arg Leu Phe Pro Gln Lys Asp Ile Tyr Leu Val Glu Tyr
35 40 45
Ser Leu Ser Thr Ser Pro Phe Val Arg Asn Arg Tyr Thr His Val Lys
50 55 60
Asp Glu Val Arg Tyr Glu Val Asn Cys Ser Gly Ile Tyr Glu Gln Glu
65 70 75 80
Pro Leu Glu Ile Gly Lys Ser Leu Glu Ile Arg Arg Arg Asp Ile Ile
85 90 95
Asp Leu Glu Asp Asp Asp Val Val Ala Met Thr Ser Asp Cys Asp Ile
100 105 110
Tyr Gln Thr Leu Lys Gly Tyr Ala *
115 120

<210> 1545
<211> 70
<212> PRT
<213> Homo sapiens

<400> 1545
Met Phe Leu Leu Lys Trp Pro Leu Trp Val Leu Gln Tyr Val Val Cys
1 5 10 15
Ser Leu Lys Asp Lys Ile His Lys Phe Phe Tyr Ile Glu Arg Val Val
20 25 30
Gly Glu Leu Arg Val Leu Pro Gln Gly Trp Met Val Ala Leu Ile Leu
35 40 45
Arg Lys Asp Phe Val Leu Pro Ser Pro Ser Asp Val Val Asn Ala Ser
50 55 60

```
Gln Pro Gly Gln Val  *
 65             69


     <210> 1546
     <211> 58
     <212> PRT
     <213> Homo sapiens


     <400> 1546
Met Tyr Gly Met Leu Glu Trp Pro Ile Ser Met Tyr Phe Val Ala Phe
  1               5                  10                  15
Leu His Cys Phe Leu Cys Ser Gly Gly Asn Leu Gly Asp Ser Phe Gln
             20                  25                  30
Ala Leu Pro Glu Leu Cys Ala Asn Cys Ser Ser Ser Pro Arg Val Leu
         35                  40                  45
Cys Cys Val Val Met Ser Pro Leu Pro  *
     50                  55      57


     <210> 1547
     <211> 65
     <212> PRT
     <213> Homo sapiens


     <400> 1547
Met Trp Leu His Glu Asn Leu Gln Phe Leu Leu Gln Leu Ile Phe His
  1               5                  10                  15
Phe Tyr Trp Thr Val Pro Pro Trp Arg Asp Trp Cys Lys Val Ile Gln
             20                  25                  30
Gln Ala Arg Asp Arg Pro Gly Pro Asn Pro Leu Leu Pro Leu Arg Met
         35                  40                  45
Gly Ala Trp His Leu Pro Gly His Asp Gly Leu Gly Arg Val Cys Thr
     50                  55                  60                  64
 *


     <210> 1548
     <211> 78
     <212> PRT
     <213> Homo sapiens


     <400> 1548
Met Phe Ile Ile Phe Leu Ala Phe Ile Ala Leu Lys Arg Ser Lys Ser
  1               5                  10                  15
Val Ile Gly Ala Phe Leu Tyr Leu Ala Ser Ile Phe Leu Ala His Gly
             20                  25                  30
Val Ala Ala His Ile Val Phe Met Ser Ala Phe Tyr Gln Ala Cys Arg
         35                  40                  45
Thr Tyr Leu Trp Trp Ala Leu Cys Glu Asn Leu Arg Met Lys Ser Val
     50                  55                  60
Ser Cys Met Leu Leu Lys Gly Met Ala Cys Leu Leu Thr  *
```

65 70 75 77

<210> 1549
<211> 54
<212> PRT
<213> Homo sapiens

<400> 1549
Met Leu Tyr Ile Glu Cys Lys Ser His Lys Leu Val Ala Pro Leu Ala
1 5 10 15
Val Phe Phe Ala Leu Phe Phe Leu Leu Ile Phe Phe Trp Val Ala Phe
20 25 30
Ser Tyr Pro Phe Glu Leu Leu Phe Leu Gln Leu Arg Ser Arg Gln Ala
35 40 45
Asp Ile Gly Val Gln *
50 53

<210> 1550
<211> 70
<212> PRT
<213> Homo sapiens

<400> 1550
Met Val Asn Thr Trp Leu Ala Ala Cys Cys Thr Val Val Thr Trp Phe
1 5 10 15
Pro Lys Met Ser Met Leu Pro Leu Pro Pro Ser Lys Pro Ser Ala Arg
20 25 30
Ser Ser Leu Trp Ile Gly Ala Pro Leu Ala Ser Arg Leu Ala Ser Thr
35 40 45
Thr Ser Leu Pro Leu Trp Cys Leu Val Glu Thr Trp Pro Arg Tyr Arg
50 55 60
Glu Leu Cys Ala Cys *
65 69

<210> 1551
<211> 224
<212> PRT
<213> Homo sapiens

<400> 1551
Met Arg Gln Ile Asn Lys Lys Gly Phe Trp Ser Tyr Gly Pro Val Ile
1 5 10 15
Leu Val Val Leu Val Val Ala Val Val Ala Ser Ser Val Asn Ser Tyr
20 25 30
Tyr Ser Ser Pro Ala Gln Gln Val Pro Lys Asn Pro Ala Leu Glu Ala
35 40 45
Phe Leu Ala Gln Phe Ser Gln Leu Glu Asp Lys Phe Pro Gly Gln Ser
50 55 60
Ser Phe Leu Trp Gln Arg Gly Arg Lys Phe Leu Gln Lys His Leu Asn
65 70 75 80

```
Ala Ser Asn Pro Thr Glu Pro Ala Thr Ile Ile Phe Thr Ala Ala Arg
                85                  90                  95
Glu Gly Arg Glu Thr Leu Lys Cys Leu Ser His His Val Ala Asp Ala
            100                 105                 110
Tyr Thr Ser Ser Gln Lys Val Ser Pro Ile Gln Ile Asp Gly Ala Gly
        115                 120                 125
Arg Thr Trp Gln Asp Ser Asp Thr Val Lys Leu Leu Val Asp Leu Glu
    130                 135                 140
Leu Ser Tyr Gly Phe Glu Asn Gly Gln Lys Ala Ala Val Val His His
145                 150                 155                 160
Phe Glu Ser Phe Pro Ala Gly Ser Thr Leu Ile Phe Tyr Lys Tyr Cys
                165                 170                 175
Asp His Glu Asn Ala Ala Phe Lys Asp Val Ala Leu Val Leu Thr Val
            180                 185                 190
Leu Leu Glu Glu Glu Thr Leu Glu Ala Ser Val Gly Pro Arg Glu Thr
        195                 200                 205
Glu Glu Lys Val Arg Asp Leu Leu Trp Ala Lys Phe Thr Asn Ser  *
    210                 215                 220         223


     <210> 1552
     <211> 57
     <212> PRT
     <213> Homo sapiens


     <400> 1552
Met Arg Gln Lys Phe Leu Lys Pro Leu Leu Ile Leu Leu His Arg Leu
  1               5                  10                  15
Lys Leu Gly Ser Leu Tyr Thr Pro Ser Ser Val Ala Arg Tyr Asp Ser
            20                  25                  30
Ser Val Asn Glu Asn Arg Ser Val Asn Ser Ser Ala Tyr Glu Glu Ala
        35                  40                  45
Lys Glu Leu Met Leu Ser Met Asn  *
    50                  55   56


     <210> 1553
     <211> 241
     <212> PRT
     <213> Homo sapiens


     <400> 1553
Met Ser Cys Val Leu Gly Gly Val Ile Pro Leu Gly Leu Leu Phe Leu
  1               5                  10                  15
Val Cys Gly Ser Gln Gly Tyr Leu Leu Pro Asn Val Thr Leu Leu Glu
            20                  25                  30
Glu Leu Leu Ser Lys Tyr Gln His Asn Glu Ser His Ser Arg Val Arg
        35                  40                  45
Arg Ala Ile Pro Arg Glu Asp Lys Glu Glu Ile Leu Met Leu His Asn
    50                  55                  60
Lys Leu Arg Gly Gln Val Gln Pro Gln Ala Ser Asn Met Glu Tyr Met
65                  70                  75                  80
Thr Trp Asp Asp Glu Leu Glu Lys Ser Ala Ala Ala Trp Ala Ser Gln
                85                  90                  95
Cys Ile Trp Glu His Gly Pro Thr Ser Leu Leu Val Ser Ile Gly Gln
```

```
            100                     105                     110
Asn Leu Gly Ala His Trp Gly Arg Tyr Arg Ser Pro Gly Phe His Val
        115                     120                     125
Gln Ser Trp Tyr Asp Glu Val Lys Asp Tyr Thr Tyr Pro Tyr Pro Ser
    130                     135                     140
Glu Cys Asn Pro Trp Cys Pro Glu Arg Cys Ser Gly Pro Met Cys Thr
145                     150                     155                 160
His Tyr Thr Gln Ile Val Trp Ala Thr Thr Asn Lys Ile Gly Cys Ala
                165                     170                     175
Val Asn Thr Cys Arg Lys Met Thr Val Trp Gly Glu Val Trp Glu Asn
            180                     185                     190
Ala Val Tyr Phe Val Cys Asn Tyr Ser Pro Lys Gly Asn Trp Ile Gly
        195                     200                     205
Glu Ala Pro Tyr Lys Asn Gly Arg Pro Cys Ser Glu Cys Pro Pro Ser
    210                     215                     220
Tyr Gly Gly Ser Cys Arg Asn Asn Leu Cys Tyr Arg Glu Glu Thr Tyr
225                     230                     235                 240
Thr
241
```

```
     <210> 1554
     <211> 56
     <212> PRT
     <213> Homo sapiens


     <400> 1554
Met Leu Thr Ser Ser Gly Cys Glu Lys His Leu Ser Leu Ala Ser Val
  1               5                  10                  15
Ser Ser Leu Ser Leu Phe Cys Val Cys Cys Ser Ser Cys Gln Leu Leu
             20                  25                  30
Trp Glu Asn Glu Cys Glu Arg Gly Ser Gln Arg Gly Trp Pro Pro Gln
        35                  40                  45
Cys Lys Trp Gly Ser Ala Val  *
    50                  55
```

```
     <210> 1555
     <211> 64
     <212> PRT
     <213> Homo sapiens


     <400> 1555
Met Tyr Gly Trp Thr Met Thr Ser Thr Ile Ser Cys Val Phe Trp Ala
  1               5                  10                  15
Cys Pro Gln Arg Lys Lys Gly Leu Cys Lys Arg Glu Gly Val Gly Ser
             20                  25                  30
Ser Ile Leu Ile His Ser Leu Ala Ala Phe Val Met Phe Asp Cys Asn
        35                  40                  45
Leu Pro Leu Leu Val Arg Arg Val Arg Arg Ile His Tyr Pro Ala  *
    50                  55                  60          63
```

```
     <210> 1556
```

<211> 71
<212> PRT
<213> Homo sapiens

<400> 1556
Met Ser Arg Pro Met Met Thr Ser Ala Ser Trp Thr Ser Val Trp Ser
1               5                   10                  15
Val Phe Val Met Ile Tyr Leu Tyr Phe Glu Arg Lys Tyr Val Leu Pro
            20                  25                  30
Leu Leu Gly Val Val Phe Tyr Thr Ile Ile Ser Asn Asp Ala Phe Ala
        35                  40                  45
Leu Glu Ser Leu Leu Ser Gly Ile Ser Thr Ser Ala Phe Phe Cys Lys
    50                  55                  60
Glu Leu Met Cys Ile Leu  *
65                  70

<210> 1557
<211> 126
<212> PRT
<213> Homo sapiens

<400> 1557
Met Gln Thr His Leu Gly Ala Ser Cys Leu Ser Leu Val Ile Arg Ile
1               5                   10                  15
Ala Leu Leu Phe Leu Val Gln Arg Asp Gly His Leu His Ser Arg Arg
            20                  25                  30
Glu Ile Tyr Ala Ile Phe Thr Lys Gly Ser Leu Cys Pro Ala Phe Lys
        35                  40                  45
Trp Ala Arg Val Gly Arg Glu Leu Phe Leu His Leu Leu Leu Ser Asn
    50                  55                  60
Cys His Gln Leu Lys Ile Ile Leu Ile Pro Lys Cys His Ile Leu Gly
65                  70                  75                  80
Trp His Ile Leu Ile Pro Phe Thr Ser Lys Ile Trp Asp Ser Tyr Phe
                85                  90                  95
Ile Val Gln Cys Phe Ser His Phe Thr Thr Leu Ala Asn Val Phe Met
            100                 105                 110
Glu Glu Asp Asn Pro Val Ser Glu Leu Gln Val Phe Gln  *
        115                 120                 125

<210> 1558
<211> 135
<212> PRT
<213> Homo sapiens

<400> 1558
Met Lys Gly Ser Ile Phe Thr Leu Phe Leu Phe Ser Val Leu Phe Ala
1               5                   10                  15
Ile Ser Glu Val Arg Ser Lys Glu Ser Val Arg Leu Cys Gly Leu Glu
            20                  25                  30
Tyr Ile Arg Thr Val Ile Tyr Ile Cys Ala Ser Ser Arg Trp Arg Arg
        35                  40                  45
His Leu Glu Gly Ile Pro Gln Ala Gln Gln Ala Glu Thr Gly Asn Ser

```
    50                  55                  60
Phe Gln Leu Pro His Lys Arg Glu Phe Ser Glu Glu Asn Pro Ala Gln
65                  70                  75                  80
Asn Leu Pro Lys Val Asp Ala Ser Gly Glu Asp Arg Leu Trp Gly Gly
                85                  90                  95
Gln Met Pro Thr Glu Glu Leu Trp Lys Ser Lys Lys His Ser Val Met
            100                 105                 110
Ser Arg Gln Asp Leu Gln Thr Leu Cys Cys Thr Asp Gly Cys Ser Met
        115                 120                 125
Thr Asp Leu Ser Ala Leu Cys
    130                 135
```

<210> 1559
<211> 203
<212> PRT
<213> Homo sapiens

<400> 1559

```
Met Glu Leu Trp Gly Ala Tyr Leu Leu Leu Cys Leu Phe Ser Leu Leu
1               5                   10                  15
Thr Gln Val Thr Thr Glu Pro Pro Thr Gln Lys Pro Lys Lys Ile Val
            20                  25                  30
Asn Ala Lys Lys Asp Val Val Asn Thr Lys Met Phe Glu Glu Leu Lys
        35                  40                  45
Ser Arg Leu Asp Thr Leu Ala Gln Glu Val Ala Leu Leu Lys Glu Gln
    50                  55                  60
Gln Ala Leu Gln Thr Val Cys Leu Lys Gly Thr Lys Val His Met Lys
65                  70                  75                  80
Cys Phe Leu Ala Phe Thr Gln Thr Lys Thr Phe His Glu Ala Ser Glu
                85                  90                  95
Asp Cys Ile Ser Arg Gly Gly Thr Leu Ser Thr Pro Gln Thr Gly Ser
            100                 105                 110
Glu Asn Asp Ala Leu Tyr Glu Tyr Leu Arg Gln Ser Val Gly Asn Glu
        115                 120                 125
Ala Glu Ile Trp Leu Gly Leu Asn Asp Met Ala Ala Glu Gly Thr Trp
    130                 135                 140
Val Asp Met Thr Gly Ala Arg Ile Ala Tyr Lys Asn Trp Glu Thr Glu
145                 150                 155                 160
Ile Thr Ala Gln Pro Asp Gly Gly Lys Thr Glu Asn Cys Ala Val Leu
                165                 170                 175
Ser Gly Ala Ala Asn Gly Lys Trp Phe Asp Lys Arg Cys Arg Asp Gln
            180                 185                 190
Leu Pro Tyr Ile Cys Gln Phe Gly Ile Val  *
        195                 200         202
```

<210> 1560
<211> 59
<212> PRT
<213> Homo sapiens

<400> 1560

```
Met Met Gly Val Ser Gly Cys Met Val Leu Leu Ala Pro Leu Leu Ala
1               5                   10                  15
```

Arg Arg Ser Gln Ser Ser Leu Trp Lys Gln Phe Glu Lys Cys Ser Ala
20 25 30
Gly Pro Lys Leu Met Leu Ser Lys Phe Leu Pro Trp Gly Lys Leu Ala
35 40 45
Met Pro Ser Arg Met Ser Asn Phe Ser Pro *
50 55 58

<210> 1561
<211> 50
<212> PRT
<213> Homo sapiens

<400> 1561
Met Lys Phe Ser Asn Val Leu Cys Thr Cys Leu Leu Ile Leu Gln Lys
1 5 10 15
Val Lys Leu Phe Tyr Lys Thr Val His Glu Asn Ser Ser Phe Leu Pro
20 25 30
Cys Phe Ser His Leu Ile Pro Ser Pro Gln Arg Asn Leu Ser Ser Ile
35 40 45
Phe *
49

<210> 1562
<211> 49
<212> PRT
<213> Homo sapiens

<400> 1562
Met Leu Phe Ser Ala Val Lys Leu Tyr Cys Cys Gln Phe Trp His Leu
1 5 10 15
Ile Leu Asn Arg Val Pro Ser Pro Ser Leu Leu Tyr Ser Cys Gly Leu
20 25 30
Ser Thr Asn Val Leu Asn Thr Thr Val Cys Tyr Val Arg Asp Lys Lys
35 40 45 48
*

<210> 1563
<211> 69
<212> PRT
<213> Homo sapiens

<400> 1563
Met Glu Arg Leu Arg Gly Lys Cys Leu Leu Ile Ile Ala Leu Met Thr
1 5 10 15
Pro Leu Cys Thr Thr Thr Ile Ser Ser Ser Cys Ile Glu Gly Ser Ala
20 25 30
Asn Phe Phe Cys Lys Glu Pro Gly Ser Asn Cys Val Phe Glu Ala Leu
35 40 45
Trp Ala Ile Trp Ser Val Gly Gln Leu Leu Ser Ser Ser Val Val Ala

```
    50                  55                  60
His Lys Gln Pro  *
 65         68


     <210> 1564
     <211> 53
     <212> PRT
     <213> Homo sapiens


     <400> 1564
Met Gln Arg Leu Gly Lys Ala Pro Gly Thr Trp Gln Ala Ile Ser Lys
  1               5                  10                  15
Cys Trp Leu Leu Leu Leu Leu Ser Leu Pro Phe Ser Gln Ser Ile Ile
             20                  25                  30
Ile Ser Leu Arg Ala Gly Thr Met Ser Tyr Leu Pro Leu Tyr Phe Pro
         35                  40                  45
Gln Tyr Phe Pro  *
    50      52


     <210> 1565
     <211> 236
     <212> PRT
     <213> Homo sapiens


     <400> 1565
Met Pro Arg Arg Gly Leu Ile Leu His Thr Arg Thr His Trp Leu Leu
  1               5                  10                  15
Leu Gly Leu Ala Leu Leu Cys Ser Leu Val Leu Phe Met Tyr Leu Leu
             20                  25                  30
Glu Cys Ala Pro Gln Thr Asp Gly Asn Ala Ser Leu Pro Gly Val Val
         35                  40                  45
Gly Glu Asn Tyr Gly Lys Glu Tyr Tyr Gln Ala Leu Leu Gln Glu Gln
    50                  55                  60
Glu Glu His Tyr Gln Thr Arg Ala Thr Ser Leu Lys Arg Gln Ile Ala
 65                  70                  75                  80
Gln Leu Lys Gln Glu Leu Gln Glu Met Ser Glu Lys Met Arg Ser Leu
                 85                  90                  95
Gln Glu Arg Arg Asn Val Gly Ala Asn Gly Ile Gly Tyr Gln Ser Asn
            100                 105                 110
Lys Glu Gln Ala Pro Ser Asp Leu Leu Glu Phe Leu His Ser Gln Ile
        115                 120                 125
Asp Lys Ala Glu Val Ser Ile Gly Ala Lys Leu Pro Ser Glu Tyr Gly
    130                 135                 140
Val Ile Pro Phe Glu Ser Phe Thr Leu Met Lys Val Phe Gln Leu Glu
145                 150                 155                 160
Met Gly Leu Thr Arg His Pro Glu Glu Lys Pro Val Arg Lys Asp Lys
                165                 170                 175
Arg Asp Glu Leu Val Glu Val Ile Glu Ala Gly Leu Glu Val Ile Asn
            180                 185                 190
Asn Pro Asp Glu Asp Asp Glu Gln Glu Asp Glu Glu Gly Pro Leu Gly
        195                 200                 205
Glu Lys Leu Ile Phe Asn Glu Asn Asp Phe Val Glu Gly Tyr Tyr Arg
    210                 215                 220
```

```
Thr Glu Arg Asp Lys Gly Thr Gln Tyr Glu Leu Phe
225                 230                 235 236


      &lt;210&gt; 1566
      &lt;211&gt; 77
      &lt;212&gt; PRT
      &lt;213&gt; Homo sapiens


      &lt;400&gt; 1566
Met Thr Ala Gly Ile Met Pro Leu Gly Leu Cys Pro Cys Ser Cys Leu
  1               5                  10                  15
Cys Leu His Ser Arg Thr Gly Ala Phe Ser Ala Val His Trp Ser Pro
             20                  25                  30
Val Glu Gly Thr Pro Asp Pro Ser Leu Arg Glu Val Ile Ser Lys Gly
         35                  40                  45
Cys Phe Ile Thr Val Phe Pro Gln Asn Asp Pro Ile Asp Thr Val Phe
     50                  55                  60
Ser Gln Cys Pro Leu Thr Phe Glu His Ile Arg Glu  *
 65                  70                  75  76


      &lt;210&gt; 1567
      &lt;211&gt; 104
      &lt;212&gt; PRT
      &lt;213&gt; Homo sapiens


      &lt;400&gt; 1567
Met Leu Ile Gly Leu Leu Ala Trp Leu Gln Thr Val Pro Ala His Gly
  1               5                  10                  15
Cys Gln Phe Leu Pro Ile Thr Ser Val Thr Ala Thr Val Tyr His Leu
             20                  25                  30
Pro Val His Gln Leu Lys Gly Arg Ser Arg Val Gln Lys Asn Leu Thr
         35                  40                  45
Leu Asp Asn Glu Gly Glu Gly Thr Trp Thr Thr Cys Leu Glu Phe Leu
     50                  55                  60
Glu Ser Leu Ala Gly Trp Arg Leu Gly Trp Gly Val Ser Arg Gly Val
 65                  70                  75                  80
Arg Glu Trp Leu Cys Leu Gln Gln Val Ser Leu His Gln Thr Pro Gly
                 85                  90                  95
Leu Pro His Lys Gln Asp Leu  *
            100         103


      &lt;210&gt; 1568
      &lt;211&gt; 46
      &lt;212&gt; PRT
      &lt;213&gt; Homo sapiens


      &lt;400&gt; 1568
Met Val Val Asn Thr Met Ile Tyr Phe Phe Ile Phe Thr Tyr Thr Leu
  1               5                  10                  15
Ala Lys Arg Ala Arg Val His Ile Asn Lys Asn Gly Asn Lys Ala Leu
```

```
             20                 25                  30
Ala Glu Lys Asn Met His Leu Thr Asn His Val Asn Ser  *
        35                  40                  45




     <210> 1569
     <211> 50
     <212> PRT
     <213> Homo sapiens


     <400> 1569
Met Leu Met Met Asp Thr Leu Trp Pro Ile Leu Leu Gln Thr Leu Lys
  1               5                  10                  15
Val Ile Ser Gln Val Gly His Ala Gly Pro Leu Ala Asn Met Ile His
             20                  25                  30
Asp Asn Pro Cys Ile Ile Ala Tyr Arg Ile Thr Leu Arg Leu Val Gly
        35                  40                  45
Pro  *
 49




     <210> 1570
     <211> 50
     <212> PRT
     <213> Homo sapiens


     <400> 1570
Met Val Gly Phe Asp Leu Leu Pro Leu Leu Phe Phe Pro Phe Phe Phe
  1               5                  10                  15
Pro Ser Leu Ile Phe Phe Pro Phe Phe Ser Ser Pro Ser Pro Ser Phe
             20                  25                  30
Gln Phe Leu Pro His Gln Glu Lys Ser Gln His Val Phe Pro Pro Asn
        35                  40                  45
Ala  *
 49




     <210> 1571
     <211> 50
     <212> PRT
     <213> Homo sapiens


     <400> 1571
Met Tyr Leu Trp Val Val Arg Trp Lys Trp Cys Leu Gln Lys Leu Gly
  1               5                  10                  15
Arg Arg Ile Leu Leu His Ser Leu His Asp Val Phe Ile Ala Asn Met
             20                  25                  30
Asp Asp Lys Gly Leu Cys Tyr Arg Gly Leu Arg Ala Pro Ser Phe Leu
        35                  40                  45
Leu  *
 49
```

<210> 1572
<211> 80
<212> PRT
<213> Homo sapiens

<400> 1572
Met Ser Ser Gly Arg Asn Phe Gly Phe Cys Phe Gln Trp Leu Pro Trp
1 5 10 15
Ala Leu Val Ala Thr Trp Ala Ser Val Thr Val Leu Met Ser Ser His
20 25 30
Ser Ser Ser Val Gly Ser Gly Leu Cys Pro Met Asp Phe Cys Ser Ser
35 40 45
Ser Arg Arg Leu Phe Ser Arg Phe Ser Ser Ile Ser Phe Leu Leu Ala
50 55 60
Ser Leu Leu Leu Ser Ser Ser Thr Lys Ser Val Ala Met Pro Thr *
65 70 75 79

<210> 1573
<211> 52
<212> PRT
<213> Homo sapiens

<400> 1573
Met Ile Asp Ile Val Arg Phe Ala Gly Leu Pro Ser Leu Leu Leu His
1 5 10 15
Ala Leu Cys Leu Ile Ser Leu Thr Tyr Pro Ser Ser Phe Arg His Ser
20 25 30
Ser Tyr Leu Ile Ser Pro Cys Ala Ser Phe Trp Ile Leu Tyr Leu Phe
35 40 45
Arg Pro Val *
50 51

<210> 1574
<211> 200
<212> PRT
<213> Homo sapiens

<400> 1574
Met Arg Leu Ser Leu Pro Leu Leu Leu Leu Leu Leu Gly Ala Trp Ala
1 5 10 15
Ile Pro Gly Gly Leu Gly Val Met Ala Pro Leu Thr Ala Thr Ala Pro
20 25 30
Glu Val Asp Asp Glu Glu Met Tyr Ser Ala His Met Pro Ala His Leu
35 40 45
Arg Cys Asp Ala Cys Arg Ala Val Ala Tyr Gln Glu Cys Gly Pro Lys
50 55 60
Thr Leu Ala Lys Ala Glu Thr Lys Leu His Thr Ser Asn Ser Gly Gly
65 70 75 80
Arg Arg Asp Val Ser Glu Leu Val Tyr Thr Asp Val Leu Asp Arg Ser
85 90 95
Cys Ser Arg Asn Trp Gln Asp Tyr Gly Val Arg Glu Val Asp Gln Val

```
            100                 105                 110
Lys Arg Leu Thr Gly Pro Gly Leu Ser Glu Gly Pro Glu Pro Ser Ile
        115                 120                 125
Ser Val Met Val Thr Gly Gly Pro Trp His Thr Arg Leu Ser Arg Thr
    130                 135                 140
Cys Leu His Tyr Leu Gly Glu Phe Gly Glu Asp Gln Ile Tyr Glu Ala
145                 150                 155                 160
His Gln Gln Gly Arg Gly Ala Leu Glu Ala Leu Leu Cys Gly Gly Pro
                165                 170                 175
Pro Gly Gly Leu Leu Arg Glu Gly Val Ser His Lys Arg Arg Ala Leu
            180                 185                 190
Val Leu Asp Ser Thr Leu Leu  *
        195             199
```

```
<210> 1575
<211> 51
<212> PRT
<213> Homo sapiens

<221> misc_feature
<222> (1)...(51)
<223> Xaa = any amino acid or nothing


<400> 1575
Met Leu Leu Gly Phe Gly Asn Val Phe Ile Leu Leu Ile Leu Xaa Thr
 1               5                  10                  15
Ala Ile Leu Trp Leu Lys Gly Ser Gln Arg Val Pro Glu Glu Pro Gly
            20                  25                  30
Glu Gln Pro Ile Tyr Met Asn Phe Ser Glu Pro Leu Thr Lys Asp Met
        35                  40                  45
Ala Thr  *
    50
```

```
<210> 1576
<211> 124
<212> PRT
<213> Homo sapiens


<400> 1576
Met Arg Ile Arg Leu Leu Cys Cys Val Ala Phe Ser Leu Leu Trp Ala
 1               5                  10                  15
Gly Pro Val Ile Ala Gly Ile Thr Gln Ala Pro Thr Ser Gln Ile Leu
            20                  25                  30
Ala Ala Gly Arg Arg Met Thr Leu Arg Cys Thr Gln Asp Met Arg His
        35                  40                  45
Asn Ala Met Tyr Trp Tyr Arg Gln Asp Leu Gly Leu Gly Leu Arg Leu
    50                  55                  60
Ile His Tyr Ser Asn Thr Ala Gly Thr Thr Gly Lys Gly Glu Val Pro
65                  70                  75                  80
Asp Gly Tyr Ser Val Ser Arg Ala Asn Thr Asp Asp Phe Pro Leu Thr
                85                  90                  95
Leu Ala Ser Ala Val Pro Ser Gln Thr Ser Val Tyr Phe Cys Ala Ser
            100                 105                 110
```

Ser Asp Gly Ala Ser Gly Ser Pro His Thr Gly Glu
115 120 124

<210> 1577
<211> 860
<212> PRT
<213> Homo sapiens

<400> 1577
Met Ala Cys Arg Trp Ser Thr Lys Glu Ser Pro Arg Trp Arg Ser Ala
1 5 10 15
Leu Leu Leu Leu Phe Leu Ala Gly Val Tyr Gly Asn Gly Ala Leu Ala
20 25 30
Glu His Ser Glu Asn Val His Ile Ser Gly Val Ser Thr Ala Cys Gly
35 40 45
Glu Thr Pro Glu Gln Ile Arg Ala Pro Ser Gly Ile Ile Thr Ser Pro
50 55 60
Gly Trp Pro Ser Glu Tyr Pro Ala Lys Ile Asn Cys Ser Trp Phe Ile
65 70 75 80
Arg Ala Asn Pro Gly Glu Ile Ile Thr Ile Ser Phe Gln Asp Phe Asp
85 90 95
Ile Gln Gly Ser Arg Arg Cys Asn Leu Asp Trp Leu Thr Ile Glu Thr
100 105 110
Tyr Lys Asn Ile Glu Ser Tyr Arg Ala Cys Gly Ser Thr Ile Pro Pro
115 120 125
Pro Tyr Ile Ser Ser Gln Asp His Ile Trp Ile Arg Phe His Ser Asp
130 135 140
Asp Asn Ile Ser Arg Lys Gly Phe Arg Leu Ala Tyr Phe Ser Gly Lys
145 150 155 160
Ser Glu Glu Pro Asn Cys Ala Cys Asp Gln Phe Arg Cys Gly Asn Gly
165 170 175
Lys Cys Ile Pro Glu Ala Trp Lys Cys Asn Asn Met Asp Glu Cys Gly
180 185 190
Asp Arg Ser Asp Glu Glu Ile Cys Ala Lys Glu Ala Asn Pro Pro Thr
195 200 205
Ala Ala Ala Phe Gln Pro Cys Ala Tyr Asn Gln Phe Gln Cys Leu Ser
210 215 220
Arg Phe Thr Lys Val Tyr Thr Cys Leu Pro Glu Ser Leu Lys Cys Asp
225 230 235 240
Gly Asn Ile Asp Cys Leu Asp Leu Gly Asp Glu Ile Asp Cys Asp Val
245 250 255
Pro Thr Cys Gly Gln Trp Leu Lys Tyr Phe Tyr Gly Thr Phe Asn Ser
260 265 270
Pro Asn Tyr Pro Asp Phe Tyr Pro Pro Gly Ser Asn Cys Thr Trp Leu
275 280 285
Ile Asp Thr Gly Asp His Arg Lys Val Ile Leu Arg Phe Thr Asp Phe
290 295 300
Lys Leu Asp Gly Thr Gly Tyr Gly Asp Tyr Val Lys Ile Tyr Asp Gly
305 310 315 320
Leu Glu Glu Asn Pro His Lys Leu Leu Arg Val Leu Thr Ala Phe Asp
325 330 335
Ser His Ala Pro Leu Thr Val Val Ser Ser Ser Gly Gln Ile Arg Val
340 345 350
His Phe Cys Ala Asp Lys Val Asn Ala Ala Arg Gly Phe Asn Ala Thr
355 360 365
Tyr Gln Val Asp Gly Phe Cys Leu Pro Trp Glu Ile Pro Cys Gly Gly

```
        370                 375                 380
Asn Trp Gly Cys Tyr Thr Glu Gln Gln Arg Cys Asp Gly Tyr Trp His
385                 390                 395                 400
Cys Pro Asn Gly Arg Asp Glu Thr Asn Cys Thr Met Cys Gln Lys Glu
                405                 410                 415
Glu Phe Pro Cys Ser Arg Asn Gly Val Cys Tyr Pro Arg Ser Asp Arg
            420                 425                 430
Cys Asn Tyr Gln Asn His Cys Pro Asn Gly Ser Asp Glu Lys Asn Cys
        435                 440                 445
Phe Phe Cys Gln Pro Gly Asn Phe His Cys Lys Asn Asn Arg Cys Val
    450                 455                 460
Phe Glu Ser Trp Val Cys Asp Ser Gln Asp Asp Cys Gly Asp Gly Ser
465                 470                 475                 480
Asp Glu Glu Asn Cys Pro Val Ile Val Pro Thr Arg Val Ile Thr Ala
                485                 490                 495
Ala Val Ile Gly Ser Leu Ile Cys Gly Leu Leu Leu Val Ile Ala Leu
            500                 505                 510
Gly Cys Thr Cys Lys Leu Tyr Ser Leu Arg Met Phe Glu Arg Arg Ser
        515                 520                 525
Phe Glu Thr Gln Leu Ser Arg Val Glu Ala Glu Leu Leu Arg Arg Glu
    530                 535                 540
Ala Pro Pro Ser Tyr Gly Gln Leu Ile Ala Gln Gly Leu Ile Pro Pro
545                 550                 555                 560
Val Glu Asp Phe Pro Val Cys Ser Pro Asn Gln Ala Ser Val Leu Glu
                565                 570                 575
Asn Leu Arg Leu Ala Val Arg Ser Gln Leu Gly Phe Thr Ser Val Arg
            580                 585                 590
Leu Pro Met Ala Gly Arg Ser Ser Asn Ile Trp Asn Arg Ile Phe Asn
        595                 600                 605
Phe Ala Arg Ser Arg His Ser Gly Ser Leu Ala Leu Val Ser Ala Asp
    610                 615                 620
Gly Asp Glu Val Val Pro Ser Gln Ser Thr Ser Arg Glu Pro Glu Arg
625                 630                 635                 640
Asn His Thr His Arg Ser Leu Phe Ser Val Glu Ser Asp Asp Thr Asp
                645                 650                 655
Thr Glu Asn Glu Arg Arg Asp Met Ala Gly Ala Ser Gly Gly Val Ala
            660                 665                 670
Ala Pro Leu Pro Gln Lys Val Pro Pro Thr Thr Ala Val Glu Ala Thr
        675                 680                 685
Val Gly Ala Cys Ala Ser Ser Ser Thr Gln Ser Thr Arg Gly Gly His
    690                 695                 700
Ala Asp Asn Gly Arg Asp Val Thr Ser Val Glu Pro Pro Ser Val Ser
705                 710                 715                 720
Pro Ala Arg His Gln Leu Thr Ser Ala Leu Ser Arg Met Thr Gln Gly
                725                 730                 735
Leu Arg Trp Val Arg Phe Thr Leu Gly Arg Ser Ser Ser Leu Ser Gln
            740                 745                 750
Asn Gln Ser Pro Leu Arg Gln Leu Asp Asn Gly Val Ser Gly Arg Glu
        755                 760                 765
Asp Asp Asp Asp Val Glu Met Leu Ile Pro Ile Ser Asp Gly Ser Ser
    770                 775                 780
Asp Phe Asp Val Asn Asp Cys Ser Arg Pro Leu Leu Asp Leu Ala Ser
785                 790                 795                 800
Asp Gln Gly Gln Gly Leu Arg Gln Pro Tyr Asn Ala Thr Asn Pro Gly
                805                 810                 815
Val Arg Pro Ser Asn Arg Asp Gly Pro Cys Glu Arg Cys Gly Ile Val
            820                 825                 830
His Thr Ala Gln Ile Pro Asp Thr Cys Leu Glu Val Thr Leu Lys Asn
        835                 840                 845
```

Glu Thr Ser Asp Asp Glu Ala Leu Leu Leu Cys *
850 855 859

<210> 1578
<211> 58
<212> PRT
<213> Homo sapiens

<400> 1578
Met Tyr Gly Met Leu Glu Trp Pro Ile Ser Met Tyr Phe Val Ala Phe
1 5 10 15
Leu His Cys Phe Leu Cys Ser Gly Gly Asn Leu Gly Asp Ser Phe Gln
20 25 30
Ala Leu Pro Glu Leu Cys Ala Asn Cys Ser Ser Ser Pro Arg Val Leu
35 40 45
Cys Cys Val Val Met Ser Pro Leu Pro *
50 55 57

<210> 1579
<211> 572
<212> PRT
<213> Homo sapiens

<400> 1579
Met Arg Arg Arg Ser Arg Met Leu Leu Cys Phe Ala Phe Leu Trp Val
1 5 10 15
Leu Gly Ile Ala Tyr Tyr Met Tyr Ser Gly Gly Gly Ser Ala Leu Ala
20 25 30
Gly Gly Ala Gly Gly Gly Ala Gly Arg Lys Glu Asp Trp Asn Glu Ile
35 40 45
Asp Pro Ile Lys Lys Lys Asp Leu His His Ser Asn Gly Glu Glu Lys
50 55 60
Ala Gln Ser Met Glu Thr Leu Pro Pro Gly Lys Val Arg Trp Pro Asp
65 70 75 80
Phe Asn Gln Glu Ala Tyr Val Gly Gly Thr Met Val Arg Ser Gly Gln
85 90 95
Asp Pro Tyr Ala Arg Asn Lys Phe Asn Gln Val Glu Ser Asp Lys Leu
100 105 110
Arg Met Asp Arg Ala Ile Pro Asp Thr Arg His Asp Gln Cys Gln Arg
115 120 125
Lys Gln Trp Arg Val Asp Leu Pro Ala Thr Ser Val Val Ile Thr Phe
130 135 140
His Asn Glu Ala Arg Ser Ala Leu Leu Arg Thr Val Val Ser Val Leu
145 150 155 160
Lys Lys Ser Pro Pro His Leu Ile Lys Glu Ile Ile Leu Val Asp Asp
165 170 175
Tyr Ser Asn Asp Pro Glu Asp Gly Ala Leu Leu Gly Lys Ile Glu Lys
180 185 190
Val Arg Val Leu Arg Asn Asp Arg Arg Glu Gly Leu Met Arg Ser Arg
195 200 205
Val Arg Gly Ala Asp Ala Ala Gln Ala Lys Val Leu Thr Phe Leu Asp
210 215 220
Ser His Cys Glu Cys Asn Glu His Trp Leu Glu Pro Leu Leu Glu Arg

225 230 235 240
Val Ala Glu Asp Arg Thr Arg Val Val Ser Pro Ile Ile Asp Val Ile
245 250 255
Asn Met Asp Asn Phe Gln Tyr Val Gly Ala Ser Ala Asp Leu Lys Gly
260 265 270
Gly Phe Asp Trp Asn Leu Val Phe Lys Trp Asp Tyr Met Thr Pro Glu
275 280 285
Gln Arg Arg Ser Arg Gln Gly Asn Pro Val Ala Pro Ile Lys Thr Pro
290 295 300
Met Ile Ala Gly Gly Leu Phe Val Met Asp Lys Phe Tyr Phe Glu Glu
305 310 315 320
Leu Gly Lys Tyr Asp Met Met Met Asp Val Trp Gly Gly Glu Asn Leu
325 330 335
Glu Ile Ser Phe Arg Val Trp Gln Cys Gly Gly Ser Leu Glu Ile Ile
340 345 350
Pro Cys Ser Arg Val Gly His Val Phe Arg Lys Gln His Pro Tyr Thr
355 360 365
Phe Pro Gly Gly Ser Gly Thr Val Phe Ala Arg Asn Thr Arg Arg Ala
370 375 380
Ala Glu Val Trp Met Asp Glu Tyr Lys Asn Phe Tyr Tyr Ala Ala Val
385 390 395 400
Pro Ser Ala Arg Asn Val Pro Tyr Gly Asn Ile Gln Ser Arg Leu Glu
405 410 415
Leu Arg Lys Lys Leu Ser Cys Lys Pro Phe Lys Trp Tyr Leu Glu Asn
420 425 430
Val Tyr Pro Glu Leu Arg Val Pro Asp His Gln Asp Ile Ala Phe Gly
435 440 445
Ala Leu Gln Gln Gly Thr Asn Cys Leu Asp Thr Leu Gly His Phe Ala
450 455 460
Asp Gly Val Val Gly Val Tyr Glu Cys His Asn Ala Gly Gly Asn Gln
465 470 475 480
Glu Trp Ala Leu Thr Lys Glu Lys Ser Val Lys His Met Asp Leu Cys
485 490 495
Leu Thr Val Val Asp Arg Ala Pro Gly Ser Leu Ile Lys Leu Gln Gly
500 505 510
Cys Arg Glu Asn Asp Ser Arg Gln Lys Trp Glu Gln Ile Glu Gly Asn
515 520 525
Ser Lys Leu Arg His Val Gly Ser Asn Leu Cys Leu Asp Ser Arg Thr
530 535 540
Ala Lys Ser Gly Gly Leu Ser Val Glu Val Cys Gly Pro Ala Leu Ser
545 550 555 560
Gln Gln Trp Lys Phe Thr Leu Asn Leu Gln Gln *
565 570 571

<210> 1580
<211> 77
<212> PRT
<213> Homo sapiens

<400> 1580
Met Glu Arg Pro Leu Cys Ser His Leu Cys Ser Cys Leu Ala Met Leu
1 5 10 15
Ala Leu Leu Ser Pro Leu Ser Leu Ala Gln Tyr Asp Ser Trp Pro His
20 25 30
Tyr Pro Glu Tyr Phe Gln Gln Pro Ala Pro Glu Tyr His Gln Pro Gln
35 40 45

Ala Pro Ala Asn Val Ala Lys Ile Gln Leu Arg Leu Ala Gly Gln Lys
50 55 60
Arg Lys His Ser Glu Gly Pro Gly Gly Gly Val Leu *
65 70 75 76

<210> 1581
<211> 494
<212> PRT
<213> Homo sapiens

<400> 1581
Met Gly Ser Leu Gln Pro Leu Ala Thr Leu Tyr Leu Leu Gly Met Leu
1 5 10 15
Val Ala Ser Cys Leu Gly Arg Leu Ser Trp Tyr Asp Pro Asp Phe Gln
20 25 30
Ala Arg Leu Thr Arg Ser Asn Ser Lys Cys Gln Gly Gln Leu Glu Val
35 40 45
Tyr Leu Lys Asp Gly Trp His Met Val Cys Ser Gln Ser Trp Gly Arg
50 55 60
Ser Ser Lys Gln Trp Glu Asp Pro Ser Gln Ala Ser Lys Val Cys Gln
65 70 75 80
Arg Leu Asn Cys Gly Val Pro Leu Ser Leu Gly Pro Phe Leu Val Thr
85 90 95
Tyr Thr Pro Gln Ser Ser Ile Ile Cys Tyr Gly Gln Leu Gly Ser Phe
100 105 110
Ser Asn Cys Ser His Ser Arg Asn Asp Met Cys His Ser Leu Gly Leu
115 120 125
Thr Cys Leu Glu Pro Gln Lys Thr Thr Pro Pro Thr Thr Arg Pro Pro
130 135 140
Pro Thr Thr Thr Pro Glu Pro Thr Ala Pro Pro Arg Leu Gln Leu Val
145 150 155 160
Ala Gln Ser Gly Gly Gln His Cys Ala Gly Val Val Glu Phe Tyr Ser
165 170 175
Gly Ser Leu Gly Gly Thr Ile Ser Tyr Glu Ala Gln Asp Lys Thr Gln
180 185 190
Asp Leu Glu Asn Phe Leu Cys Asn Asn Leu Gln Cys Gly Ser Phe Leu
195 200 205
Lys His Leu Pro Glu Thr Glu Ala Gly Arg Ala Gln Asp Pro Gly Glu
210 215 220
Pro Arg Glu His Gln Pro Leu Pro Ile Gln Trp Lys Ile Gln Asn Ser
225 230 235 240
Ser Cys Thr Ser Leu Glu His Cys Phe Arg Lys Ile Lys Pro Gln Lys
245 250 255
Ser Gly Arg Val Leu Ala Leu Leu Cys Ser Gly Phe Gln Pro Lys Val
260 265 270
Gln Ser Arg Leu Val Gly Gly Ser Ser Ile Cys Glu Gly Thr Val Glu
275 280 285
Val Arg Gln Gly Ala Gln Trp Ala Ala Leu Cys Asp Ser Ser Ser Ala
290 295 300
Arg Ser Ser Leu Arg Trp Glu Glu Val Cys Arg Glu Gln Gln Cys Gly
305 310 315 320
Ser Val Asn Ser Tyr Arg Val Leu Asp Ala Gly Asp Pro Thr Ser Arg
325 330 335
Gly Leu Phe Cys Pro His Gln Lys Leu Ser Gln Cys His Glu Leu Trp
340 345 350
Glu Arg Asn Ser Tyr Cys Lys Lys Val Phe Val Thr Cys Gln Asp Pro

355 360 365
Asn Pro Ala Gly Leu Ala Ala Gly Thr Val Ala Ser Ile Ile Leu Ala
370 375 380
Leu Val Leu Leu Val Val Leu Leu Val Val Cys Gly Pro Leu Ala Tyr
385 390 395 400
Lys Lys Leu Val Lys Lys Phe Arg Gln Lys Lys Gln Arg Gln Trp Ile
405 410 415
Gly Pro Thr Gly Met Asn Gln Asn Met Ser Phe His Arg Asn His Thr
420 425 430
Ala Thr Val Arg Ser His Ala Glu Asn Pro Thr Ala Ser His Val Asp
435 440 445
Asn Glu Tyr Ser Gln Pro Pro Arg Asn Ser Arg Leu Ser Ala Tyr Pro
450 455 460
Ala Leu Glu Gly Ala Leu His Arg Ser Ser Met Gln Pro Asp Asn Ser
465 470 475 480
Ser Asp Ser Asp Tyr Asp Leu His Gly Ala Gln Arg Leu *
485 490 493

<210> 1582
<211> 329
<212> PRT
<213> Homo sapiens

<400> 1582
Met Gln Gly Leu Cys Ile Ser Val Ala Val Phe Leu His Tyr Phe Leu
1 5 10 15
Leu Val Ser Phe Thr Trp Met Gly Leu Glu Ala Phe His Met Tyr Leu
20 25 30
Ala Leu Val Lys Val Phe Asn Thr Tyr Ile Arg Lys Tyr Ile Leu Lys
35 40 45
Phe Cys Ile Val Gly Trp Gly Val Pro Ala Val Val Val Thr Ile Ile
50 55 60
Leu Thr Ile Ser Pro Asp Asn Tyr Gly Leu Gly Ser Tyr Gly Lys Phe
65 70 75 80
Pro Asn Gly Ser Pro Asp Asp Phe Cys Trp Ile Asn Asn Asn Ala Val
85 90 95
Phe Tyr Ile Thr Val Val Gly Tyr Phe Cys Val Ile Phe Leu Leu Asn
100 105 110
Val Ser Met Phe Ile Val Val Leu Val Gln Leu Cys Arg Ile Lys Lys
115 120 125
Lys Lys Gln Leu Gly Ala Gln Arg Lys Thr Ser Ile Gln Asp Leu Arg
130 135 140
Ser Ile Ala Gly Leu Thr Phe Leu Leu Gly Ile Thr Trp Gly Phe Ala
145 150 155 160
Phe Phe Ala Trp Gly Pro Val Asn Val Thr Phe Met Tyr Leu Phe Ala
165 170 175
Ile Phe Asn Thr Leu Gln Gly Phe Phe Ile Phe Ile Phe Tyr Cys Val
180 185 190
Ala Lys Glu Asn Val Arg Lys Gln Trp Arg Arg Tyr Leu Cys Cys Gly
195 200 205
Lys Leu Arg Leu Ala Glu Asn Ser Asp Trp Ser Lys Thr Ala Thr Asn
210 215 220
Gly Leu Lys Lys Gln Thr Val Asn Gln Gly Val Ser Ser Ser Ser Asn
225 230 235 240
Ser Leu Gln Ser Ser Ser Asn Ser Thr Asn Ser Thr Thr Leu Leu Val
245 250 255

```
Asn Asn Asp Cys Ser Val His Ala Ser Gly Asn Gly Asn Ala Ser Thr
            260                 265                 270
Glu Arg Asn Gly Val Ser Phe Ser Val Gln Asn Gly Asp Val Cys Leu
        275                 280                 285
His Asp Phe Thr Gly Lys Gln His Met Phe Asn Glu Lys Glu Asp Ser
    290                 295                 300
Cys Asn Gly Lys Gly Arg Met Ala Leu Arg Arg Thr Ser Lys Arg Gly
305                 310                 315                 320
Ser Leu His Phe Ile Glu Gln Met  *
                325         328



     <210> 1583
     <211> 49
     <212> PRT
     <213> Homo sapiens


     <400> 1583
Met Gly Met Gly Arg Leu Leu Pro Met Ala Trp Val Leu Ala Gly Ile
 1               5                  10                  15
Pro Thr Gly Ala Gln Gln Ser Trp Arg Arg Pro Trp Ser Gly Ser Ala
            20                  25                  30
Pro Arg Cys Ala Ser Cys Gly Ser Ala Trp Arg Cys Cys Ala Val Arg
        35                  40                  45              48
 *



     <210> 1584
     <211> 671
     <212> PRT
     <213> Homo sapiens


     <400> 1584
Met Ile Ala Ser Cys Leu Cys Tyr Leu Leu Leu Pro Ala Thr Arg Leu
 1               5                  10                  15
Phe Arg Ala Leu Ser Asp Ala Phe Phe Thr Cys Arg Lys Asn Val Leu
            20                  25                  30
Leu Ala Asn Ser Ser Ser Pro Gln Val Glu Gly Asp Phe Ala Met Ala
        35                  40                  45
Pro Arg Gly Pro Glu Gln Glu Glu Cys Glu Gly Leu Leu Gln Gln Trp
    50                  55                  60
Arg Glu Glu Gly Leu Ser Gln Val Leu Ser Thr Ala Ser Glu Gly Pro
65                  70                  75                  80
Leu Ile Asp Lys Gly Leu Ala Gln Ser Ser Leu Ala Leu Leu Met Asp
                85                  90                  95
Asn Pro Gly Glu Glu Asn Ala Ala Ser Glu Asp Arg Trp Ser Ser Arg
            100                 105                 110
Gln Leu Ser Asp Leu Arg Ala Ala Glu Asn Leu Asp Glu Pro Phe Pro
        115                 120                 125
Glu Met Leu Gly Glu Glu Pro Leu Leu Glu Val Glu Gly Val Glu Gly
    130                 135                 140
Ser Met Trp Ala Ala Ile Pro Met Gln Ser Glu Pro Gln Tyr Ala Asp
145                 150                 155                 160
Cys Ala Ala Leu Pro Val Gly Ala Leu Ala Thr Glu Gln Trp Glu Glu
```

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                165                  170                  175
Asp Pro Ala Val Leu Ala Trp Ser Ile Ala Pro Glu Pro Val Pro Gln
            180                  185                  190
Glu Glu Ala Ser Ile Trp Pro Phe Glu Gly Leu Gly Gln Leu Gln Pro
        195                  200                  205
Pro Ala Val Glu Ile Pro Tyr His Glu Ile Leu Trp Arg Glu Trp Glu
    210                  215                  220
Asp Phe Ser Thr Gln Pro Asp Ala Gln Gly Leu Lys Ala Gly Asp Gly
225                  230                  235                  240
Pro Gln Phe Gln Phe Thr Leu Met Ser Tyr Asn Ile Leu Ala Gln Asp
                245                  250                  255
Leu Met Gln Gln Ser Ser Glu Leu Tyr Leu His Cys His Pro Asp Ile
            260                  265                  270
Leu Asn Trp Asn Tyr Arg Phe Val Asn Leu Met Gln Glu Phe Gln His
        275                  280                  285
Trp Asp Pro Asp Ile Leu Cys Leu Gln Glu Val Gln Glu Asp His Tyr
    290                  295                  300
Trp Glu Gln Leu Glu Pro Ser Leu Arg Met Met Gly Phe Thr Cys Phe
305                  310                  315                  320
Tyr Lys Arg Arg Thr Gly Cys Lys Thr Asp Gly Cys Ala Val Cys Tyr
                325                  330                  335
Lys Pro Thr Arg Phe Arg Leu Leu Cys Ala Ser Pro Val Glu Tyr Phe
            340                  345                  350
Arg Pro Gly Leu Glu Leu Leu Asn Arg Asp Asn Val Gly Leu Val Leu
        355                  360                  365
Leu Leu Gln Pro Leu Val Pro Glu Gly Leu Gly Gln Val Ser Val Ala
    370                  375                  380
Pro Leu Cys Val Ala Asn Thr His Ile Leu Tyr Asn Pro Arg Arg Gly
385                  390                  395                  400
Asp Val Lys Leu Ala Gln Met Ala Ile Leu Leu Ala Glu Val Asp Lys
                405                  410                  415
Val Ala Arg Leu Ser Asp Gly Ser His Cys Pro Ile Ile Leu Cys Gly
            420                  425                  430
Asp Leu Asn Ser Val Pro Asp Ser Pro Leu Tyr Asn Phe Ile Arg Asp
        435                  440                  445
Gly Glu Leu Gln Tyr His Gly Met Pro Ala Trp Lys Val Ser Gly Gln
    450                  455                  460
Glu Asp Phe Ser His Gln Leu Tyr Gln Arg Lys Leu Gln Ala Pro Leu
465                  470                  475                  480
Trp Pro Ser Ser Leu Gly Ile Thr Asp Cys Cys Gln Tyr Val Thr Ser
                485                  490                  495
Cys His Pro Lys Arg Ser Glu Arg Arg Lys Tyr Gly Arg Asp Phe Leu
            500                  505                  510
Leu Arg Phe Arg Phe Cys Ser Ile Ala Cys Gln Arg Pro Val Gly Leu
        515                  520                  525
Val Leu Met Glu Gly Val Thr Asp Thr Lys Pro Glu Arg Pro Ala Gly
    530                  535                  540
Trp Ala Glu Ser Val Leu Glu Glu Asp Ala Ser Glu Leu Glu Pro Ala
545                  550                  555                  560
Phe Ser Arg Thr Val Gly Thr Ile Gln His Cys Leu His Leu Thr Ser
                565                  570                  575
Val Tyr Thr His Phe Leu Pro Gln Arg Gly Arg Pro Glu Val Thr Thr
            580                  585                  590
Met Pro Leu Gly Leu Gly Met Thr Val Asp Tyr Ile Phe Phe Ser Ala
        595                  600                  605
Glu Ser Cys Glu Asn Gly Asn Arg Thr Asp His Arg Leu Tyr Arg Asp
    610                  615                  620
Gly Thr Leu Lys Leu Leu Gly Arg Leu Ser Leu Leu Ser Glu Glu Ile
625                  630                  635                  640
```

Leu Trp Ala Ala Asn Gly Leu Pro Asn Pro Phe Cys Ser Ser Asp His
645 650 655
Leu Cys Leu Leu Ala Ser Leu Gly Met Glu Val Thr Ala Pro *
660 665 670

<210> 1585
<211> 318
<212> PRT
<213> Homo sapiens

<400> 1585
Met Met Cys Leu Lys Ile Leu Arg Ile Ser Leu Ala Ile Leu Ala Gly
1 5 10 15
Trp Ala Leu Cys Ser Ala Asn Ser Glu Leu Gly Trp Thr Arg Lys Lys
20 25 30
Ser Leu Val Glu Arg Glu His Leu Asn Gln Val Leu Leu Glu Gly Glu
35 40 45
Arg Cys Trp Leu Gly Ala Lys Val Arg Arg Pro Arg Ala Ser Pro Gln
50 55 60
His His Leu Phe Gly Val Tyr Pro Ser Arg Ala Gly Asn Tyr Leu Arg
65 70 75 80
Pro Tyr Pro Val Gly Glu Gln Glu Ile His His Thr Gly Arg Ser Lys
85 90 95
Pro Asp Thr Glu Gly Asn Ala Val Ser Leu Val Pro Pro Asp Leu Thr
100 105 110
Glu Asn Pro Ala Gly Leu Arg Gly Ala Val Glu Glu Pro Ala Ala Pro
115 120 125
Trp Val Gly Asp Ser Pro Ile Gly Gln Ser Glu Leu Leu Gly Asp Asp
130 135 140
Asp Ala Tyr Leu Gly Asn Gln Arg Ser Lys Glu Ser Leu Gly Glu Ala
145 150 155 160
Gly Ile Gln Lys Gly Ser Ala Met Ala Ala Thr Thr Thr Thr Ala Ile
165 170 175
Phe Thr Thr Leu Asn Glu Pro Lys Pro Glu Thr Gln Arg Arg Gly Trp
180 185 190
Ala Lys Ser Arg Gln Arg Arg Gln Val Trp Lys Arg Arg Ala Glu Asp
195 200 205
Gly Gln Gly Asp Ser Gly Ile Ser Ser His Phe Gln Pro Trp Pro Lys
210 215 220
His Ser Leu Lys His Arg Val Lys Lys Ser Pro Pro Glu Glu Ser Asn
225 230 235 240
Gln Asn Gly Gly Glu Gly Ser Tyr Arg Glu Ala Glu Thr Phe Asn Ser
245 250 255
Gln Val Gly Leu Pro Ile Leu Tyr Phe Ser Gly Arg Arg Glu Arg Leu
260 265 270
Leu Leu Arg Pro Glu Val Leu Ala Glu Ile Pro Arg Glu Ala Phe Thr
275 280 285
Val Glu Ala Trp Val Lys Pro Glu Gly Gly Gln Asn Asn Pro Ala Ile
290 295 300
Ile Ala Gly Asn Thr Leu Leu Leu Gly Phe Leu Lys Ser *
305 310 315 317

<210> 1586
<211> 80

<212> PRT
<213> Homo sapiens

<400> 1586
Met Ile Ala Leu Thr Gln Leu Leu Thr Phe Ile Leu Ser Cys Asn Ser
1 5 10 15
Ser Leu Leu His Ile Phe Pro Phe Cys Glu Gln Val Leu Val Glu Asn
20 25 30
Gly Thr Lys Ala Gly His Ser Leu Leu Met Asp Ala Arg Asp Leu Val
35 40 45
Leu Lys Gly Lys Glu Lys Ser Pro Leu Asp Pro Arg Pro Gly Phe Val
50 55 60
Phe Ala Pro Val Ser Ile Thr Ser Ala Cys Pro Thr Thr Arg Ile *
65 70 75 79

<210> 1587
<211> 316
<212> PRT
<213> Homo sapiens

<400> 1587
Met Phe Phe Gly Ser Ala Ala Leu Gly Thr Leu Thr Gly Leu Ile Ser
1 5 10 15
Ala Leu Val Leu Lys His Ile Asp Leu Arg Lys Thr Pro Ser Leu Glu
20 25 30
Phe Gly Met Met Ile Ile Phe Ala Tyr Leu Pro Tyr Gly Leu Ala Glu
35 40 45
Gly Ile Ser Leu Ser Gly Ile Met Ala Ile Leu Phe Ser Gly Ile Val
50 55 60
Met Ser His Tyr Thr His His Asn Leu Ser Pro Val Thr Gln Ile Leu
65 70 75 80
Met Gln Gln Thr Leu Arg Thr Val Ala Phe Leu Cys Glu Thr Cys Val
85 90 95
Phe Ala Phe Leu Gly Leu Ser Ile Phe Ser Phe Pro His Lys Phe Glu
100 105 110
Ile Ser Phe Val Ile Trp Cys Ile Val Leu Val Leu Phe Gly Arg Ala
115 120 125
Val Asn Ile Phe Pro Leu Ser Tyr Leu Leu Asn Phe Phe Arg Asp His
130 135 140
Lys Ile Thr Pro Lys Met Met Phe Ile Met Trp Phe Ser Gly Leu Arg
145 150 155 160
Gly Ala Ile Pro Tyr Ala Leu Ser Leu His Leu Asp Leu Glu Pro Met
165 170 175
Glu Lys Arg Gln Leu Ile Gly Thr Thr Thr Ile Val Ile Val Leu Phe
180 185 190
Thr Ile Leu Leu Leu Gly Gly Ser Thr Met Pro Leu Ile Arg Leu Met
195 200 205
Asp Ile Glu Asp Ala Lys Ala His Arg Arg Asn Lys Lys Asp Val Asn
210 215 220
Leu Ser Lys Thr Glu Lys Met Gly Asn Thr Val Glu Ser Glu His Leu
225 230 235 240
Ser Glu Leu Thr Glu Glu Glu Tyr Glu Ala His Tyr Ile Arg Arg Gln
245 250 255
Asp Leu Lys Gly Phe Val Trp Leu Asp Ala Lys Tyr Leu Asn Pro Phe
260 265 270

Phe Thr Arg Arg Leu Thr Gln Glu Asp Leu His His Gly Arg Ile Gln
275 280 285
Met Lys Thr Leu Thr Asn Lys Trp Tyr Glu Glu Val Arg Gln Gly Pro
290 295 300
Ser Gly Ser Glu Asp Asp Glu Gln Glu Leu Leu *
305 310 315

<210> 1588
<211> 53
<212> PRT
<213> Homo sapiens

<221> misc_feature
<222> (1)...(53)
<223> Xaa = any amino acid or nothing

<400> 1588
Met Cys Ser Leu Met Phe Gly Ser Ser Val Phe Val Cys Phe Pro Pro
1 5 10 15
Cys Val Pro Leu Pro Ala Pro His Ser Gly Gly Pro Pro His Arg Ala
20 25 30
Gly Arg Ser Val Phe Ser Ala Met Lys Leu Gly Lys Xaa Arg Ser His
35 40 45
Lys Glu Glu Pro Gln
50 53

<210> 1589
<211> 437
<212> PRT
<213> Homo sapiens

<400> 1589
Met Leu Lys Val Ser Ala Val Leu Cys Val Cys Ala Ala Ala Trp Cys
1 5 10 15
Ser Gln Ser Leu Ala Ala Ala Ala Ala Val Ala Ala Ala Gly Gly Arg
20 25 30
Ser Asp Gly Gly Asn Phe Leu Asp Asp Lys Gln Trp Leu Thr Thr Ile
35 40 45
Ser Gln Tyr Asp Lys Glu Val Gly Gln Trp Asn Lys Phe Arg Asp Glu
50 55 60
Val Glu Asp Asp Tyr Phe Arg Thr Trp Ser Pro Gly Lys Pro Phe Asp
65 70 75 80
Gln Ala Leu Asp Pro Ala Lys Asp Pro Cys Leu Lys Met Lys Cys Ser
85 90 95
Arg His Lys Val Cys Ile Ala Gln Asp Ser Gln Thr Ala Val Cys Ile
100 105 110
Ser His Arg Arg Leu Thr His Arg Met Lys Glu Ala Gly Val Asp His
115 120 125
Arg Gln Trp Arg Gly Pro Ile Leu Ser Thr Cys Lys Gln Cys Pro Val
130 135 140
Val Tyr Pro Ser Pro Val Cys Gly Ser Asp Gly His Thr Tyr Ser Phe
145 150 155 160
Gln Cys Lys Leu Glu Tyr Gln Ala Cys Val Leu Gly Lys Gln Ile Ser

```
                    165                     170                     175
Val Lys Cys Glu Gly His Cys Pro Cys Pro Ser Asp Lys Pro Thr Ser
                180                     185                     190
Thr Ser Arg Asn Val Lys Arg Ala Cys Ser Asp Leu Glu Phe Arg Glu
            195                     200                     205
Val Ala Asn Arg Leu Arg Asp Trp Phe Lys Ala Leu His Glu Ser Gly
        210                     215                     220
Ser Gln Asn Lys Lys Thr Lys Thr Leu Leu Arg Pro Glu Arg Ser Arg
225                     230                     235                     240
Phe Asp Thr Ser Ile Leu Pro Ile Cys Lys Asp Ser Leu Gly Trp Met
                    245                     250                     255
Phe Asn Arg Leu Asp Thr Asn Tyr Asp Leu Leu Leu Asp Gln Ser Glu
                260                     265                     270
Leu Arg Ser Ile Tyr Leu Asp Lys Asn Glu Gln Cys Thr Lys Ala Phe
            275                     280                     285
Phe Asn Ser Cys Asp Thr Tyr Lys Asp Ser Leu Ile Ser Asn Asn Glu
        290                     295                     300
Trp Cys Tyr Cys Phe Gln Arg Gln Gln Asp Pro Pro Cys Gln Thr Glu
305                     310                     315                     320
Leu Ser Asn Ile Gln Lys Arg Gln Gly Val Lys Lys Leu Leu Gly Gln
                    325                     330                     335
Tyr Ile Pro Leu Cys Asp Glu Asp Gly Tyr Tyr Lys Pro Thr Gln Cys
                340                     345                     350
His Gly Ser Val Gly Gln Cys Trp Cys Val Asp Arg Tyr Gly Asn Glu
            355                     360                     365
Val Met Gly Ser Arg Ile Asn Gly Val Ala Asp Cys Ala Ile Asp Phe
        370                     375                     380
Glu Ile Ser Gly Asp Phe Ala Ser Gly Asp Phe His Glu Trp Thr Asp
385                     390                     395                     400
Asp Glu Asp Asp Glu Asp Asp Ile Met Asn Asp Glu Asp Glu Ile Glu
                    405                     410                     415
Asp Asp Asp Glu Asp Glu Gly Asp Asp Asp Asp Gly Gly Asp Asp His
                420                     425                     430
Asp Val Tyr Ile  *
            435 436
```

```
<210> 1590
<211> 49
<212> PRT
<213> Homo sapiens
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```
<400> 1590
Met Phe Gln Ile Tyr Phe Ser Phe Cys Gln Leu Cys Phe Ile Trp Ser
 1               5                  10                  15
Cys Phe Phe Asn Ser Arg Glu Thr Phe Asn Glu Ile Tyr Lys Phe Phe
                20                  25                  30
Leu Lys Ser Val Met Val Arg Lys Ile Phe Glu Cys His Lys Met Ser
            35                  40                  45              48
 *
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<210> 1591
<211> 73
<212> PRT
```

<213> Homo sapiens

<400> 1591
Met Ser Leu Asn Val Leu Leu Ala Leu Phe Cys Leu Leu Leu Ala Lys
1 5 10 15
Glu Arg Thr Thr Thr Lys Arg Cys Ile Ser Cys Leu Pro Phe Ser Thr
20 25 30
Phe Phe Ser Phe Gly Pro Leu Gln Lys Val Thr Asp Pro Ser Ser Trp
35 40 45
Ala Leu Ala Phe Ser Val Cys Gln Ala Cys Thr Arg Ser Glu Leu Pro
50 55 60
Gly Ala Leu Arg Thr Arg Gly Ser Thr
65 70 73

<210> 1592
<211> 62
<212> PRT
<213> Homo sapiens

<400> 1592
Met Tyr Phe Ser Leu Ile Phe Leu Val Phe Phe Phe Leu Ser Leu Pro
1 5 10 15
Leu Ser Ser Ser Ser Ser Glu Pro Thr Ser Ser Ile Leu Gly Phe Ser
20 25 30
Ser Ser Ser Leu Ser Ser Ser Ser Phe Ser Pro Phe Ser Ser Ser Ala
35 40 45
Ser Ser Ser Leu Ile Ser Phe Ser Arg Ser Phe Ser Lys *
50 55 60 61

<210> 1593
<211> 128
<212> PRT
<213> Homo sapiens

<400> 1593
Met Arg Ala Met Leu Gly Thr Cys Ala Leu Gly Gln Phe Phe Leu Ile
1 5 10 15
Met Gly Asn Thr Gln Arg Cys Asp Asp Phe Pro Thr Glu Ser Pro Pro
20 25 30
Ala Lys Thr Asn Val Ser Arg Ala Gly Leu Ser Pro Pro Cys Glu Ala
35 40 45
Leu His Gly Val Glu Ser Arg Gly Ser Cys Ser His Gly Lys Leu Gln
50 55 60
Ser Pro Pro Gly Arg Asp Trp Pro Gln Gly Asp Pro Gln Asp Arg Pro
65 70 75 80
Lys Arg Arg Trp Gln Arg Pro Gly Pro Ala Gly Arg Gly Ala Pro Asp
85 90 95
Pro Thr Pro Lys Gly Gln Gly Ala Ala Val Pro Pro Arg Ser Ala Ser
100 105 110
Met Phe Leu Ile His Lys Gln Met Trp Ala Tyr Gly Phe Gly Asp *
115 120 125 127

<210> 1594
<211> 46
<212> PRT
<213> Homo sapiens

<400> 1594
Met Ile Trp Ala Leu Ser Ser Ser Leu Ile Pro Phe Leu Ile Ala Leu
1 5 10 15
Cys Phe Val His Ser Ala Asn Ser His Leu Gln Val Leu Val Ile Cys
20 25 30
Ser Ser Leu Phe Leu Glu Pro Pro Pro His Asn Phe Met *
35 40 45

<210> 1595
<211> 86
<212> PRT
<213> Homo sapiens

<400> 1595
Met Trp Glu Glu Leu Leu Arg Gly Leu Thr Ala Pro Tyr Trp Leu Ser
1 5 10 15
Ser Trp Leu Cys Phe Ser Trp Arg Ala Ala Thr Val Ala Val Ala Val
20 25 30
Ala Val Ala Val Ala Ala Ala Ala Ala Thr Ala Ala Ala Ala Ala Ala
35 40 45
Cys Val Lys Ser Val Glu Gly Leu Ala Ala Cys Glu Gly Arg Pro Arg
50 55 60
Pro Pro Gly Pro Pro Ala Tyr Leu Gln Glu Thr Gln Asp Cys His Ala
65 70 75 80
Leu Cys Val Gly Ser *
85

<210> 1596
<211> 69
<212> PRT
<213> Homo sapiens

<400> 1596
Met Val Leu Ser Trp Leu Thr Leu Ile Glu Ala Leu Ala Asp Val Met
1 5 10 15
Thr Thr Asp Gly Asn Met Leu Gln Leu Phe Cys Val Glu Arg Thr Asn
20 25 30
Leu Leu Val Asn Gln Ile Arg Met Thr Leu Tyr Ala Gln Tyr Arg His
35 40 45
Val Arg Pro Phe Arg Thr Ile Met Lys Pro Ile Leu Thr Arg Glu Val
50 55 60
Gln Thr Lys Asp *
65 68

<210> 1597
<211> 56
<212> PRT
<213> Homo sapiens

<400> 1597
Met Phe Leu Leu Phe Ser Arg Ile Ser Asn Leu Met Phe Val Asn His
1 5 10 15
Lys Leu Pro Met Leu Ile Thr Glu Asn Lys Gln Val Ser Lys Glu Glu
20 25 30
Asn Lys Ala Thr His Ser His Arg Ser Ser Phe Gln Ser Ser Thr Ile
35 40 45
Ser Ser Arg Leu Asn Leu Ile *
50 55

<210> 1598
<211> 97
<212> PRT
<213> Homo sapiens

<400> 1598
Met His Glu Ser Pro Leu Ala Trp Ala Ser Val His Leu Ser Ser Leu
1 5 10 15
Pro Leu Leu Cys Thr Ala Cys Ser Ser Pro Leu Met Gly Asn Ser Val
20 25 30
Leu Cys Arg Ala Pro Ala Asp Met Gly Leu Ala Trp Met Leu Leu Leu
35 40 45
Ser Glu Pro Arg Arg Val Val Pro Gly Ile Ala Ala Gln Val Leu Thr
50 55 60
Ala Leu Arg Arg Arg Leu Leu Ser Gly Thr Leu Pro Ser Phe Pro Arg
65 70 75 80
Arg Lys Asn Pro Leu His Glu His Leu Leu Ala Phe Ile Val Arg Leu
85 90 95 96
*

<210> 1599
<211> 113
<212> PRT
<213> Homo sapiens

<400> 1599
Met Thr Val Ser Gly Thr Val Val Leu Val Ala Gly Thr Leu Cys Phe
1 5 10 15
Ala Trp Trp Ser Glu Gly Asp Ala Thr Ala Gln Pro Gly Gln Leu Ala
20 25 30
Pro Pro Thr Glu Tyr Pro Val Pro Glu Gly Pro Ser Pro Leu Leu Arg
35 40 45
Ser Val Ser Phe Val Cys Cys Gly Ala Gly Gly Leu Leu Leu Leu Ile
50 55 60
Gly Leu Leu Trp Ser Val Lys Ala Ser Ile Pro Gly Pro Pro Arg Trp

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65                  70                  75                  80
Asp Pro Tyr His Leu Ser Arg Asp Leu Tyr Tyr Leu Thr Val Glu Ser
                85                  90                  95
Ser Glu Lys Glu Ser Cys Arg Thr Pro Lys Val Val Asp Ile Pro Asp
            100                 105                 110     112
*
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<210> 1600
<211> 103
<212> PRT
<213> Homo sapiens


<400> 1600
Met Gly Ala Trp Ala Trp Val Pro Thr Pro Ser Leu Cys Leu Cys His
 1              5                   10                  15
Ser Thr Cys Leu Glu Phe Leu Leu Phe Leu Tyr Ile Leu Phe Tyr Cys
            20                  25                  30
Ile Phe Glu Thr Val Ser Leu Ser Pro Arg Leu Glu Arg Ser Gly Ala
        35                  40                  45
Ile Leu Ala Arg Cys Asn Leu Cys Leu Arg Gly Ser Ser Asp Ser Arg
    50                  55                  60
Ala Leu Ala Ser Arg Val Ala Glu Thr Thr Gly Met His His His Ala
65                  70                  75                  80
Trp Leu Ile Phe Ala Phe Leu Val Glu Thr Gly Phe His His Val Gly
                85                  90                  95
Gln Ala Gly Leu Asn Ser  *
            100     102
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<210> 1601
<211> 84
<212> PRT
<213> Homo sapiens


<400> 1601
Met Val Ala Leu Leu Cys Arg Gln Ile Ile Ser Ala Ala Phe Ser Gly
 1              5                   10                  15
Glu Gly Thr Pro Leu Cys Ser Trp Ser Ser Gly Pro Ile Leu Ser Ser
            20                  25                  30
Val Cys Leu Leu Cys Pro Leu Ala Val Leu Cys Pro Ala Lys Pro Glu
        35                  40                  45
Pro Arg Ala Phe Thr Asp Leu Arg Gly Glu Glu Val Cys Ala Asp Trp
    50                  55                  60
Phe Met Gly Gly His Gly Arg Val Glu Arg Gly Thr Met Ser Pro His
65                  70                  75                  80
Ser Gly Leu  *
        83
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<210> 1602
<211> 91
<212> PRT
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<213> Homo sapiens

<400> 1602
Met Lys Thr Leu Pro Val Leu Val Leu Ser Leu Thr Leu Leu Thr Val
1               5                   10                  15
Phe Ser Glu Thr Ser Pro Ile Leu Thr Glu Lys Gln Ala Lys Gln Leu
            20                  25                  30
Leu Arg Ser Arg Arg Gln Asp Arg Pro Ser Lys Pro Gly Phe Pro Asp
        35                  40                  45
Glu Pro Met Arg Glu Tyr Met His His Leu Leu Ala Leu Glu His Arg
    50                  55                  60
Ala Glu Glu Gln Phe Leu Glu His Trp Leu Asn Pro His Cys Lys Pro
65                  70                  75                  80
His Cys Asp Arg Asn Arg Ile His Pro Val  *
                85                  90

<210> 1603
<211> 69
<212> PRT
<213> Homo sapiens

<400> 1603
Met Lys Arg Asp Val Leu Ile Thr Glu Thr Phe Cys Ile Leu Phe Trp
1               5                   10                  15
Leu Cys Ala Phe Ser Ser Met Asn Asp Tyr Val Phe Lys Pro His Val
            20                  25                  30
Leu Tyr Ile Asp Cys Pro Leu Lys Arg Leu Asp Ser Ser Val Cys Gln
        35                  40                  45
His Ile Gly Thr Glu Tyr Asn Tyr Thr Leu Ile Ile Ser Gln Val Phe
    50                  55                  60
Ile Leu Glu Val  *
65          68

<210> 1604
<211> 83
<212> PRT
<213> Homo sapiens

<400> 1604
Met Leu Gln Pro Met Phe Phe Thr Leu Ser Thr His Leu Val Gly Leu
1               5                   10                  15
Ser Gln Ile Ser Tyr Leu Ser Phe Pro Leu Ile Ser Leu His Pro Ala
            20                  25                  30
Gln Val Val Lys Arg Gln Ser Ser Leu Pro Arg Leu Met Gln Ser Ser
        35                  40                  45
Lys Glu Ser Lys Ala Val Leu Val Glu Ile Ile Leu Arg Ala Arg Lys
    50                  55                  60
Val Val Lys Tyr Ile Ser Lys Gly Phe Leu Arg Ala Val Cys Ala Glu
65                  70                  75                  80
Met Ile  *
    82

<210> 1605
<211> 110
<212> PRT
<213> Homo sapiens

<221> misc_feature
<222> (1)...(110)
<223> Xaa = any amino acid or nothing

<400> 1605
Met Ser Thr Ile Ile Phe Gln Trp Pro Phe Met Leu Val Ser Leu His
1 5 10 15
Arg Cys Arg Lys Leu Pro Arg Ala Leu Lys Asp Trp Gln Ala Phe Leu
20 25 30
Asp Leu Lys Lys Ile Ile Asp Asp Phe Ser Glu Cys Cys Pro Leu Leu
35 40 45
Glu Tyr Met Gly Ser Lys Ala Met Met Glu Arg His Xaa Glu Arg Ile
50 55 60
Thr Thr Leu Thr Gly His Ser Leu Asp Val Gly Asn Glu Ser Phe Lys
65 70 75 80
Leu Arg Asn Ile Met Glu Ala Pro Leu Leu Xaa Tyr Lys Glu Glu Ile
85 90 95
Glu Val Glu Tyr Asp Val Met Glu Asp Cys Lys Val Ser Trp
100 105 110

<210> 1606
<211> 72
<212> PRT
<213> Homo sapiens

<400> 1606
Met Thr Ala Gly Thr Val Thr Met Leu Leu Trp His Ala Ser Asn Trp
1 5 10 15
Asp Val Gln Leu Pro Ser Gln Pro Leu Val Glu Leu Thr Pro Val Arg
20 25 30
Asp Leu Asp Thr Ser Gly Leu Thr Ala Phe Leu Ala Arg Asp Met Asn
35 40 45
Leu Leu Ser Gly Asn Val Asn Thr Met Asn Gly Glu Ser Ile Ile Ala
50 55 60
Ile Thr Met Lys Met Leu Ala *
65 70 71

<210> 1607
<211> 59
<212> PRT
<213> Homo sapiens

<400> 1607
Met Phe Thr Arg Phe Ile Gly Leu Phe Leu Lys Phe Ile Leu Met Phe
1 5 10 15

Phe Leu Leu Leu Ser Phe Ile Ser Tyr Phe Cys Leu Phe Pro Cys Ser
20 25 30
Asn Leu Pro Lys Val Ile Ala Ile Phe Asn Ile Val Leu Ile Leu Ser
35 40 45
Ile Val Phe Arg Glu Ile Thr Asp Thr Tyr *
50 55 58

<210> 1608
<211> 118
<212> PRT
<213> Homo sapiens

<400> 1608
Met Leu Val Thr Asp Thr Glu Ala Phe Trp Gln Pro Gln Pro Trp Phe
1 5 10 15
Val Val Val Leu Thr Ala Thr Gly Ala Leu Leu Leu Leu Ala Leu Gly
20 25 30
Trp Leu Leu Gly Arg Leu Leu Gln Gly Leu Ala Gln Leu Leu Gln Ala
35 40 45
Pro Ser Lys Pro Ala Gln Ala Leu Leu Leu Asn Ser Ile Gln Gly Thr
50 55 60
Glu Gly Ser Ile Glu Gly Phe Leu Glu Ala Pro Lys Met Glu Met Ser
65 70 75 80
Gln Ala Pro Ser Ser Val Met Ser Leu Gln His Phe Asp Gly Arg Thr
85 90 95
Gln Asp Ser Arg Thr Gly Arg Asp Tyr Leu Val Asn Thr His Thr Gly
100 105 110
Ala Arg Arg Trp Leu *
115 117

<210> 1609
<211> 50
<212> PRT
<213> Homo sapiens

<400> 1609
Met Val Ile Gly Ser Leu His Thr Phe Thr Leu Leu Ala Ala Ser Ser
1 5 10 15
Leu Val Asp Thr Pro Lys Gln Ile Gln Leu Leu Met Gln Asn Leu Met
20 25 30
Asn Asp Pro Arg Lys Glu Val Lys Ile Leu Ala Ile Gln Asp Leu Lys
35 40 45
Leu Leu
50

<210> 1610
<211> 50
<212> PRT
<213> Homo sapiens

<400> 1610
Met Val Leu Ile Leu Ser Pro Gly Leu Ser Ile Leu Phe Thr Lys Met
1 5 10 15
Ser Glu Thr Phe Ser Ser Ser Leu Leu Lys Leu Ser Ser Ser Ile Cys
20 25 30
Ile Phe Pro Leu Cys Ile Asn Met Ile Ile Cys Tyr Gln Lys Lys Ser
35 40 45
Gln *
49

<210> 1611
<211> 56
<212> PRT
<213> Homo sapiens

<400> 1611
Met Ser Phe Gln Ala Phe Val Phe Leu Met Ile Gly Trp Leu His Pro
1 5 10 15
Asp Pro Arg Leu Met Thr Gln Arg Ser Cys Gly Pro His Pro Glu Val
20 25 30
Asp Ser Ala Gln Glu Asp His Phe Ser His Pro Tyr Asp Ile Pro Asn
35 40 45
Gln Ser Ala Pro Pro Leu Pro *
50 55

<210> 1612
<211> 75
<212> PRT
<213> Homo sapiens

<400> 1612
Met Leu Thr Leu Ala Leu Leu Val Leu Arg Ile Cys Val Cys Glu Ala
1 5 10 15
Ala Ser Thr Phe Val Cys Pro Cys Leu Pro Trp Leu Ser Leu Leu Phe
20 25 30
Leu His Leu Leu Pro Arg Leu Phe Gln Val Gln Ile Trp Phe Leu Leu
35 40 45
Phe Leu Pro Phe Leu Leu Leu Leu Pro Ser Val Pro Glu Ile Phe Pro
50 55 60
Ala Pro Gln Ala Trp Gly Leu Gly Cys Ser *
65 70 74

<210> 1613
<211> 192
<212> PRT
<213> Homo sapiens

<400> 1613
Met Phe Thr Cys Leu Phe Leu Phe Ser Ala Val Leu Arg Ala Leu Phe
1 5 10 15

Arg Lys Ser Asp Pro Lys Arg Phe Gln Asn Ile Phe Thr Thr Ile Phe
20 25 30
Thr Leu Phe Thr Leu Leu Thr Leu Asp Asp Trp Ser Leu Ile Tyr Met
35 40 45
Asp Ser Arg Ala Gln Gly Ala Trp Tyr Ile Ile Pro Ile Leu Ile Ile
50 55 60
Tyr Ile Ile Ile Gln Tyr Phe Ile Phe Leu Asn Leu Val Ile Thr Val
65 70 75 80
Leu Val Asp Ser Phe Gln Thr Ala Leu Phe Lys Gly Leu Glu Lys Ala
85 90 95
Lys Gln Glu Arg Ala Ala Arg Ile Gln Glu Lys Leu Leu Glu Asp Ser
100 105 110
Leu Thr Glu Leu Arg Ala Ala Glu Pro Lys Glu Val Ala Ser Glu Gly
115 120 125
Thr Met Leu Lys Arg Leu Ile Glu Lys Lys Phe Gly Thr Met Thr Glu
130 135 140
Lys Gln Gln Glu Leu Leu Phe His Tyr Leu Gln Leu Val Ala Ser Val
145 150 155 160
Glu Gln Glu Gln Gln Lys Phe Arg Ser Gln Ala Ala Val Ile Asp Glu
165 170 175
Ile Val Asp Thr Thr Phe Glu Ala Gly Glu Glu Asp Phe Arg Asn *
180 185 190 191

<210> 1614
<211> 153
<212> PRT
<213> Homo sapiens

<400> 1614
Met Asp Leu Val Gln Phe Phe Val Thr Phe Phe Ser Cys Phe Leu Ser
1 5 10 15
Leu Leu Leu Val Ala Ala Val Val Trp Lys Ile Lys Gln Thr Cys Trp
20 25 30
Ala Ser Arg Arg Arg Glu Gln Leu Leu Arg Glu Arg Gln Gln Met Ala
35 40 45
Ser Arg Pro Phe Ala Ser Val Asp Val Ala Leu Glu Val Gly Ala Glu
50 55 60
Gln Thr Glu Phe Leu Arg Gly Pro Leu Glu Gly Ala Pro Lys Pro Ile
65 70 75 80
Ala Ile Glu Pro Cys Ala Gly Asn Arg Ala Ala Val Leu Thr Val Phe
85 90 95
Leu Cys Leu Pro Arg Gly Ser Ser Gly Ala Pro Pro Pro Gly Gln Ser
100 105 110
Gly Leu Ala Ile Ala Ser Ala Leu Ile Asp Ile Ser Gln Gln Lys Ala
115 120 125
Ser Asp Ser Lys Asp Lys Thr Ser Gly Val Arg Asn Arg Lys His Leu
130 135 140
Ser Thr Arg Gln Gly Thr Cys Val *
145 150 152

<210> 1615
<211> 135
<212> PRT
<213> Homo sapiens

<400> 1615
Met His Trp Leu Arg Ala Ser Ala Gly Ser Leu Leu Met Val Pro Leu
1 5 10 15
Met Thr Asp Leu His Glu Leu Ala Leu Pro Pro Ala Ser Leu Arg Thr
20 25 30
Val Val Lys Glu Asn Met Cys Val Leu Pro Phe Pro Val Lys Thr Ser
35 40 45
Gly Arg Ser Leu Thr Gly Ser Ala Trp Ser Arg Phe His Leu Pro Cys
50 55 60
His Leu Arg Pro Gly Asp Arg Leu Pro Cys His Cys Leu Gly Lys Phe
65 70 75 80
Arg Lys Arg Val Ala Lys Trp Cys Ile Arg Lys Asn Met Ala Arg Ser
85 90 95
Pro His Leu Leu Gly Gly Arg Pro Asn Ser Thr Ser Gly Pro Leu Cys
100 105 110
Asp Phe Pro Ala Pro Ser Lys Gln Val Thr Pro Leu Leu Trp Val Ser
115 120 125
Val Ser Leu Pro Ile Lys *
130 134

<210> 1616
<211> 60
<212> PRT
<213> Homo sapiens

<400> 1616
Met Leu His Gln Met Lys Phe Ile Gly His Leu Ile Phe Ile Val Val
1 5 10 15
Leu Asp Pro Asp Leu Ser Asp Met Lys Asn Asn Glu Pro Tyr Asp Tyr
20 25 30
Lys Phe Val Lys Trp Met Thr Lys His Lys Val Met Phe Ile Val Leu
35 40 45
Cys Lys Ile Leu Leu Tyr Phe Ile Val Asn Phe *
50 55 59

<210> 1617
<211> 49
<212> PRT
<213> Homo sapiens

<400> 1617
Met Pro Glu His Leu Cys Phe Glu Ile Cys Asn Thr Leu Leu Asn Phe
1 5 10 15
Phe Ser Phe Leu Leu Leu Cys Val Thr Asp His Glu Thr Thr Phe Phe
20 25 30
Asp Ser Gly Trp Lys Ala Ser Gly Ser Thr Val Thr Cys Lys Ala Gly
35 40 45 48
*

<210> 1618
<211> 95
<212> PRT
<213> Homo sapiens

<400> 1618
Met Trp Thr Val Leu Trp His Arg Phe Ser Met Val Leu Arg Leu Pro
1 5 10 15
Glu Glu Ala Ser Ala Gln Glu Gly Glu Leu Ser Leu Ser Ser Pro Pro
20 25 30
Ser Pro Glu Pro Asp Trp Thr Leu Ile Ser Pro Gln Gly Met Ala Ala
35 40 45
Leu Leu Ser Leu Ala Met Ala Thr Phe Thr Gln Glu Pro Gln Leu Cys
50 55 60
Leu Ser Cys Leu Ser Gln His Gly Ser Ile Leu Met Ser Ile Leu Lys
65 70 75 80
His Leu Leu Cys Pro Ser Phe Leu Asn Gln Leu Arg Gln Ala *
85 90 94

<210> 1619
<211> 54
<212> PRT
<213> Homo sapiens

<400> 1619
Met Ile Leu Met Leu Leu Leu Leu Ile Val Asp Leu Val Gln Leu Ala
1 5 10 15
Gly Asn Ala Val Ile Ser Ser Gly Ser Trp Asp Ser Ala Cys Thr Gly
20 25 30
Thr Pro Ser Pro Ser Thr Pro Ser Thr Trp Pro Gly Pro Thr Ser Ser
35 40 45
Ser Ala Pro Arg Phe *
50 53

<210> 1620
<211> 71
<212> PRT
<213> Homo sapiens

<400> 1620
Met Cys Cys Ser Phe Leu Leu Glu Gly Leu Ile Ser Leu Phe Ser Leu
1 5 10. 15
Gln Leu Phe Ser Val Gln Leu Val Leu Leu Phe Phe Leu Trp Ile Val
20 25 30
Ser Tyr Ser Lys Lys Gln Ile Lys Asp Thr Phe Ala Lys Thr Lys Asn
35 40 45
Thr Val Ala Arg Ile Leu Leu Ser Ile Pro Asp Leu Pro Ser Leu Thr
50 55 60
Leu Ile Thr Gln Ile Leu *
65 70

<210> 1621
<211> 90
<212> PRT
<213> Homo sapiens

<221> misc_feature
<222> (1)...(90)
<223> Xaa = any amino acid or nothing

<400> 1621
Met Asp His Lys Ser Leu Trp Ala Gly Val Glu Val Leu Leu Leu Leu
1 5 10 15
Gln Gly Gly Ser Ala Tyr Lys Leu Val Cys Tyr Phe Thr Asn Trp Ser
20 25 30
Gln Asp Arg Gln Glu Pro Gly Lys Phe Thr Pro Glu Asn Ile Asp Pro
35 40 45
Phe Leu Cys Ser His Leu Ile Tyr Ser Phe Ala Ser Ile Glu Asn Asn
50 55 60
Lys Val Ile Ile Arg Thr Pro Xaa Phe Phe Pro Leu Pro Leu Gly His
65 70 75 80
Arg Leu Gln Thr Ile Asn Pro Arg Leu *
85 89

<210> 1622
<211> 53
<212> PRT
<213> Homo sapiens

<400> 1622
Met Gln Cys Ala Ile Cys Ile Leu Leu Tyr Leu Leu Asn Lys Lys Thr
1 5 10 15
Val Trp Arg Cys Ser Arg Ile His His Asn Asn Thr Val Val Leu Thr
20 25 30
Arg Glu Ser Ser Pro Phe Leu Thr Thr Cys Thr Leu Ser Ser Val Leu
35 40 45
Leu Thr Lys Ala *
50 52

<210> 1623
<211> 978
<212> PRT
<213> Homo sapiens

<400> 1623
Met Pro Ala Arg Arg Leu Leu Leu Leu Leu Thr Leu Leu Leu Pro Gly
1 5 10 15
Leu Gly Ile Phe Gly Ser Thr Ser Thr Val Thr Leu Pro Glu Thr Leu
20 25 30
Leu Phe Val Ser Thr Leu Asp Gly Ser Leu His Ala Val Ser Lys Arg
35 40 45

Thr Gly Ser Ile Lys Trp Thr Leu Lys Glu Asp Pro Val Leu Gln Val
50 55 60
Pro Thr His Val Glu Glu Pro Ala Phe Leu Pro Asp Pro Asn Asp Gly
65 70 75 80
Ser Leu Tyr Thr Leu Gly Ser Lys Asn Asn Glu Gly Leu Thr Lys Leu
85 90 95
Pro Phe Thr Ile Pro Glu Leu Val Gln Ala Ser Pro Cys Arg Ser Ser
100 105 110
Asp Gly Ile Leu Tyr Met Gly Lys Lys Gln Asp Ile Trp Tyr Val Ile
115 120 125
Asp Leu Leu Thr Gly Glu Lys Gln Gln Thr Leu Ser Ser Ala Phe Ala
130 135 140
Asp Ser Leu Cys Pro Ser Thr Ser Leu Leu Tyr Leu Gly Arg Thr Glu
145 150 155 160
Tyr Thr Ile Thr Met Tyr Asp Thr Lys Thr Arg Glu Leu Arg Trp Asn
165 170 175
Ala Thr Tyr Phe Asp Tyr Ala Ala Ser Leu Pro Glu Asp Asp Val Asp
180 185 190
Tyr Lys Met Ser His Phe Val Ser Asn Gly Asp Gly Leu Val Val Thr
195 200 205
Val Asp Ser Glu Ser Gly Asp Val Leu Trp Ile Gln Asn Tyr Ala Ser
210 215 220
Pro Val Val Ala Phe Tyr Val Trp Gln Arg Glu Gly Leu Arg Lys Val
225 230 235 240
Met His Ile Asn Val Ala Val Glu Thr Leu Arg Tyr Leu Thr Phe Met
245 250 255
Ser Gly Glu Val Gly Arg Ile Thr Lys Trp Lys Tyr Pro Phe Pro Lys
260 265 270
Glu Thr Glu Ala Lys Ser Lys Leu Thr Pro Thr Leu Tyr Val Gly Lys
275 280 285
Tyr Ser Thr Ser Leu Tyr Ala Ser Pro Ser Met Val His Glu Gly Val
290 295 300
Ala Val Val Pro Arg Gly Ser Thr Leu Pro Leu Leu Glu Gly Pro Gln
305 310 315 320
Thr Asp Gly Val Thr Ile Gly Asp Lys Gly Glu Cys Val Ile Thr Pro
325 330 335
Ser Thr Asp Val Lys Phe Asp Pro Gly Leu Lys Ser Lys Asn Lys Leu
340 345 350
Asn Tyr Leu Arg Asn Tyr Trp Leu Leu Ile Gly His His Glu Thr Pro
355 360 365
Leu Ser Ala Ser Thr Lys Met Leu Glu Arg Phe Pro Asn Asn Leu Pro
370 375 380
Lys His Arg Glu Asn Val Ile Pro Ala Asp Ser Glu Lys Lys Ser Phe
385 390 395 400
Glu Glu Val Ile Asn Leu Val Asp Gln Thr Ser Glu Asn Ala Pro Thr
405 410 415
Thr Val Ser Arg Asp Val Glu Glu Lys Pro Ala His Ala Pro Ala Arg
420 425 430
Pro Glu Ala Pro Val Asp Ser Met Leu Lys Asp Met Ala Thr Ile Ile
435 440 445
Leu Ser Thr Phe Leu Leu Ile Gly Trp Val Ala Phe Ile Ile Thr Tyr
450 455 460
Pro Leu Ser Met His Gln Gln Gln Leu Gln His Gln Gln Phe Gln
465 470 475 480
Lys Glu Leu Glu Lys Ile Gln Leu Leu Gln Gln Gln Gln Gln Leu
485 490 495
Pro Phe His Pro Pro Gly Asp Thr Ala Gln Asp Gly Glu Leu Leu Asp
500 505 510
Thr Ser Gly Pro Tyr Ser Glu Ser Ser Gly Thr Ser Ser Pro Ser Thr

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        515                 520                 525
Ser Pro Arg Ala Ser Asn His Ser Leu Cys Ser Gly Ser Ser Ala Ser
    530                 535                 540
Lys Ala Gly Ser Ser Pro Ser Leu Glu Gln Asp Asp Gly Asp Glu Glu
545                 550                 555                 560
Thr Ser Val Val Ile Val Gly Lys Ile Ser Phe Cys Pro Lys Asp Val
                565                 570                 575
Leu Gly His Gly Ala Glu Gly Thr Ile Val Tyr Arg Gly Met Phe Asp
            580                 585                 590
Asn Arg Asp Val Ala Val Lys Arg Ile Leu Pro Glu Cys Phe Ser Phe
        595                 600                 605
Ala Asp Arg Glu Val Gln Leu Leu Arg Glu Ser Asp Glu His Pro Asn
    610                 615                 620
Val Ile Arg Tyr Phe Cys Thr Glu Lys Asp Arg Gln Phe Gln Tyr Ile
625                 630                 635                 640
Ala Ile Glu Leu Cys Ala Ala Thr Leu Gln Glu Tyr Val Glu Gln Lys
                645                 650                 655
Asp Phe Ala His Leu Gly Leu Glu Pro Ile Thr Leu Leu Gln Gln Thr
            660                 665                 670
Thr Ser Gly Leu Ala His Leu His Ser Leu Asn Ile Val His Arg Asp
        675                 680                 685
Leu Lys Pro His Asn Ile Leu Ile Ser Met Pro Asn Ala His Gly Lys
    690                 695                 700
Ile Lys Ala Met Ile Ser Asp Phe Gly Leu Trp Lys Lys Leu Ala Val
705                 710                 715                 720
Gly Arg His Ser Phe Ser Arg Arg Ser Gly Val Pro Gly Thr Glu Gly
                725                 730                 735
Trp Ile Ala Pro Glu Met Leu Ser Glu Asp Cys Lys Glu Asn Pro Thr
            740                 745                 750
Tyr Thr Val Asp Ile Phe Ser Ala Gly Cys Val Phe Tyr Tyr Val Ile
        755                 760                 765
Ser Glu Gly Ser His Pro Phe Gly Lys Ser Leu Gln Arg Gln Ala Asn
    770                 775                 780
Ile Leu Leu Gly Ala Cys Ser Leu Asp Cys Leu His Pro Glu Lys His
785                 790                 795                 800
Glu Asp Val Ile Ala Arg Glu Leu Ile Glu Lys Met Ile Ala Met Asp
                805                 810                 815
Pro Gln Lys Arg Pro Ser Ala Lys His Val Leu Lys His Pro Phe Phe
            820                 825                 830
Trp Ser Leu Glu Lys Gln Leu Gln Phe Phe Gln Asp Val Ser Asp Arg
        835                 840                 845
Ile Glu Lys Glu Ser Leu Asp Gly Pro Ile Val Lys Gln Leu Glu Arg
    850                 855                 860
Gly Gly Arg Ala Val Val Lys Met Asp Trp Arg Glu Asn Ile Thr Val
865                 870                 875                 880
Pro Leu Gln Thr Asp Leu Arg Lys Phe Arg Thr Tyr Lys Gly Gly Ser
                885                 890                 895
Val Arg Asp Leu Leu Arg Ala Met Arg Asn Lys Lys His His Tyr Arg
            900                 905                 910
Glu Leu Pro Ala Glu Val Arg Glu Thr Leu Gly Thr Leu Pro Asp Asp
        915                 920                 925
Phe Val Cys Tyr Phe Thr Ser Arg Phe Pro His Leu Leu Ala His Thr
    930                 935                 940
Tyr Arg Ala Met Glu Leu Cys Ser His Glu Arg Leu Phe Gln Pro Tyr
945                 950                 955                 960
Tyr Phe His Glu Pro Pro Glu Pro Gln Pro Pro Val Thr Pro Asp Ala
                965                 970                 975
Leu *
977
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<210> 1624
<211> 56
<212> PRT
<213> Homo sapiens

<400> 1624
Met His Ser Cys Trp Thr Phe Gln Asp Leu Ser Leu Val Gln Leu Cys
1 5 10 15
Leu Pro Leu Ser Cys Pro Gln Gln Gly Pro Val Gly Pro Gly Gly Phe
20 25 30
Leu Leu Pro Val Ser Gln Val Gly Pro Pro Lys Pro Ala Gly His Trp
35 40 45
Gln Arg Lys Leu Leu Met Pro *
50 55

<210> 1625
<211> 146
<212> PRT
<213> Homo sapiens

<400> 1625
Met Glu Leu Ala Leu Leu Cys Gly Leu Val Val Met Ala Gly Val Ile
1 5 10 15
Pro Ile Gln Gly Gly Ile Leu Asn Leu Asn Lys Met Val Lys Gln Val
20 25 30
Thr Gly Lys Met Pro Ile Leu Ser Tyr Trp Pro Tyr Gly Cys His Cys
35 40 45
Gly Leu Gly Gly Arg Gly Gln Pro Lys Asp Ala Thr Asp Trp Cys Cys
50 55 60
Gln Thr His Asp Cys Cys Tyr Asp His Leu Lys Thr Gln Gly Cys Gly
65 70 75 80
Ile Tyr Lys Asp Tyr Tyr Arg Tyr Asn Phe Ser Gln Gly Asn Ile His
85 90 95
Cys Ser Asp Lys Gly Ser Trp Cys Glu Gln Gln Leu Cys Ala Cys Asp
100 105 110
Lys Glu Val Ala Phe Cys Leu Lys Arg Asn Leu Asp Thr Tyr Gln Lys
115 120 125
Arg Leu Arg Phe Tyr Trp Arg Pro His Cys Arg Gly Gln Thr Pro Gly
130 135 140
Cys *
145

<210> 1626
<211> 385
<212> PRT
<213> Homo sapiens

<400> 1626
Met Glu Phe Gly Leu Ser Trp Leu Phe Leu Val Ala Ile Leu Lys Gly

1 5 10 15
Val Gln Cys Glu Val Gln Leu Val Glu Ser Gly Gly Gly Leu Val Gln
20 25 30
Pro Gly Gly Ser Leu Arg Leu Ser Cys Ala Ala Ser Gly Phe Thr Phe
35 40 45
Ser Ser Tyr Ala Met Ser Trp Val Arg Gln Ala Pro Gly Lys Gly Leu
50 55 60
Glu Trp Val Ser Gly Ile Gly Gly Ser Gly Ser Ser Thr Tyr Tyr Ala
65 70 75 80
Asp Ser Val Lys Gly Arg Phe Thr Ile Ser Arg Asp Asn Ser Gln Asn
85 90 95
Thr Leu Tyr Leu Gln Met Asn Ser Leu Arg Ala Glu Asp Thr Ala Val
100 105 110
Tyr Tyr Cys Ala Lys Ser His Pro Ala Tyr Tyr Tyr Gly Ser Gly Ser
115 120 125
Tyr Ser Ser His Tyr Tyr Tyr Tyr Tyr Gly Met Asp Val Trp Gly Gln
130 135 140
Gly Thr Thr Val Thr Val Ser Ser Gly Asp Gly Ser Ser Gly Gly Ser
145 150 155 160
Gly Gly Ala Ser Thr Gly Glu Ile Val Leu Thr Gln Ser Pro Gly Thr
165 170 175
Leu Ser Leu Ser Pro Gly Glu Arg Ala Thr Leu Ser Cys Arg Ala Ser
180 185 190
Gln Ser Val Ser Ser Ser Tyr Leu Ala Trp Tyr Gln Gln Lys Pro Gly
195 200 205
Gln Ala Pro Arg Leu Leu Ile Tyr Gly Ala Ser Ser Arg Ala Thr Gly
210 215 220
Ile Pro Asp Arg Phe Ser Gly Ser Gly Ser Gly Thr Asp Phe Thr Leu
225 230 235 240
Thr Ile Ser Arg Leu Glu Pro Glu Asp Phe Ala Val Tyr Tyr Cys Gln
245 250 255
Gln Tyr Gly Ser Ser Pro Thr Thr Phe Gly Gln Gly Thr Lys Val Glu
260 265 270
Ile Lys Arg Thr Val Ala Ala Pro Ser Val Phe Ile Phe Pro Pro Ser
275 280 285
Asp Glu Gln Leu Lys Ser Gly Thr Ala Ser Val Val Cys Leu Leu Asn
290 295 300
Asn Phe Tyr Pro Arg Glu Ala Lys Val Gln Trp Lys Val Asp Asn Ala
305 310 315 320
Leu Gln Ser Gly Asn Ser Gln Glu Ser Val Thr Glu Gln Asp Ser Lys
325 330 335
Asp Ser Thr Tyr Ser Leu Ser Ser Thr Leu Thr Leu Ser Lys Ala Asp
340 345 350
Tyr Glu Lys His Lys Val Tyr Ala Cys Glu Val Thr His Ser Gly Ala
355 360 365
Leu Ser Phe Ala Arg Ser Gln Arg Ser Phe Gln Pro Gly Glu Ser Val
370 375 380 384
*

<210> 1627
<211> 101
<212> PRT
<213> Homo sapiens

<400> 1627

```
Met Ile Val His Cys Thr Ile Ile Pro Leu Ser Phe Cys Val His Arg
  1               5                  10                  15
Leu Arg Ala Pro Leu Asp Ala Tyr Phe Gln Val Ser Arg Thr Gln Pro
             20                  25                  30
Asp Leu Pro Ala Thr Thr Tyr Asp Ser Glu Thr Arg Asn Pro Val Ser
         35                  40                  45
Glu Glu Leu Gln Val Ser Ser Ser Ser Asp Ser Asp Ser Asp Ser Ser
     50                  55                  60
Ala Glu Tyr Gly Gly Val Val Asp Gln Ala Glu Glu Ser Gly Ala Val
 65                  70                  75                  80
Ile Leu Glu Gly Gln Tyr Phe Thr Gln Val Trp Thr His Lys Ala Asn
                 85                  90                  95
Ile His Glu Ala  *
            100
```

<210> 1628
<211> 71
<212> PRT
<213> Homo sapiens

<400> 1628

```
Met Ile Phe Tyr Val Ile Leu Ser Ser Pro Ser Ser Arg Thr Phe Phe
  1               5                  10                  15
Lys Ile Thr Leu Ile Met Ser Leu Gly Leu Ile Ser Lys Leu Leu Ile
             20                  25                  30
Thr Ser Cys Thr Phe Asp Thr Val Thr Phe Met Met Leu Thr Asn Ile
         35                  40                  45
Thr Lys Met Lys Ile Ser Ser Gly Lys Ala Thr Gln Ser Gln Glu Phe
     50                  55                  60
Phe Ser Glu Leu Ile Leu Tyr
 65                  70  71
```

<210> 1629
<211> 112
<212> PRT
<213> Homo sapiens

<400> 1629

```
Met Ala His Tyr Lys Thr Glu Gln Asp Asp Trp Leu Ile Ile Tyr Leu
  1               5                  10                  15
Lys Tyr Leu Leu Phe Val Phe Asn Phe Phe Phe Trp Val Gly Gly Ala
             20                  25                  30
Ala Val Leu Ala Val Gly Ile Trp Thr Leu Val Glu Lys Ser Gly Tyr
         35                  40                  45
Leu Ser Val Leu Ala Ser Ser Thr Phe Ala Ala Ser Ala Tyr Ile Leu
     50                  55                  60
Ile Phe Ala Gly Val Leu Val Met Val Thr Gly Phe Leu Gly Phe Gly
 65                  70                  75                  80
Ala Ile Leu Trp Glu Arg Lys Gly Cys Leu Ser Thr Tyr Phe Cys Leu
                 85                  90                  95
Leu Leu Val Ile Phe Leu Asp Glu Leu Glu Ala Gly Val Leu Ala His
            100                 105                 110     112
```

<210> 1630
<211> 47
<212> PRT
<213> Homo sapiens

<400> 1630
Met Trp Pro Gln Leu Leu Lys Ser Phe Phe Leu Ile Pro Thr Gln Ile
1 5 10 15
His Phe Asn Leu Thr Asn Leu Pro Ser Trp Arg Arg Arg Glu Leu Arg
20 25 30
Arg Phe Val Trp Val Ser Met Pro Glu Leu Ile Gly Ala Ser *
35 40 45 46

<210> 1631
<211> 79
<212> PRT
<213> Homo sapiens

<400> 1631
Met Tyr Met Trp Ser Gly Leu Leu Gly Ser Lys Trp Thr Leu Val Tyr
1 5 10 15
Ser His Phe Leu Asn Met Ala Pro Ala Ser Phe Ser His Tyr Gln Ala
20 25 30
Ser Leu Pro Leu Leu Glu His Asp Thr Leu Ser Ser Ser Arg Val His
35 40 45
Ser Tyr Gln Cys Pro Gly Phe Phe Cys Phe Phe Pro Ser Val Leu Glu
50 55 60
Phe Ser Gln Leu Gln Lys Thr Tyr Ser Leu Cys Leu Pro Phe *
65 70 75 78

<210> 1632
<211> 48
<212> PRT
<213> Homo sapiens

<400> 1632
Met Phe Met Cys Arg Leu Leu Leu Trp Ala Thr Gly Ala Tyr Gly Phe
1 5 10 15
Leu Gly Asp Asp Val Glu Tyr Thr Ser Val Leu Pro His Gln Lys Gly
20 25 30
Lys Glu Ala Trp Val Phe Ile Cys Gln Leu Pro Phe Ile Ile Gly *
35 40 45 47

<210> 1633
<211> 58
<212> PRT

<213> Homo sapiens

<400> 1633
Met Cys Leu Arg Arg Thr Leu Leu Trp His Leu His Ile Ala Pro Leu
1 5 10 15
Val Asn Ile Leu Ser Asp Tyr Lys Pro Leu Gly Arg Trp Asn His Ala
20 25 30
Pro Ala Leu Thr Ala Gly Ala Leu His Lys Thr Thr Ile Leu Leu Pro
35 40 45
Gln Gly His Pro Lys Ala Ala Asn Pro *
50 55 57

<210> 1634
<211> 55
<212> PRT
<213> Homo sapiens

<400> 1634
Met Leu Val Phe Asn Leu Ser Leu Val Leu Ser His Ser Val Leu Glu
1 5 10 15
Phe Val Met Phe Leu Tyr Ser Leu Asp Ser Ser His Val Cys Pro Leu
20 25 30
Val Val Pro Val Thr Leu Asp Leu Ile Tyr Leu Val Tyr Leu Pro Cys
35 40 45
Gln Ser Tyr Ile Leu Ile *
50 54

<210> 1635
<211> 78
<212> PRT
<213> Homo sapiens

<400> 1635
Met Ala Val Val Gln Ala Leu Thr Pro Leu Val Ser Ala Ala Ala Thr
1 5 10 15
Ala Ser Cys Leu Thr Ser Cys Ser Trp Ser Leu Thr Phe Pro Glu His
20 25 30
Ser Val Asn Tyr Gln Ser His Pro Ser Glu Thr Gln Pro Tyr Leu Leu
35 40 45
Arg Ser Thr Lys Glu Lys His His His Trp Leu Thr Ala Lys Ala Thr
50 55 60
Cys Pro Ala Ala Gly Ala Glu Gly Leu Pro Ser Arg Gly *
65 70 75 77

<210> 1636
<211> 51
<212> PRT
<213> Homo sapiens

<400> 1636
Met Phe Cys Ser Phe Pro Leu Leu Ile Leu Gln Val Tyr Pro Thr Trp
1 5 10 15
Lys Asn Pro Asn Trp His Leu Thr Phe His Thr Ser Val Phe Ser Phe
20 25 30
Pro Lys Gly Val Arg Ser Leu Ala Arg Gly Ile Pro Asp His Leu His
35 40 45
Ser Ala *
50

<210> 1637
<211> 123
<212> PRT
<213> Homo sapiens

<400> 1637
Met Gln Gln Met Met Trp Ala Gly Leu Leu Cys Pro Gln Leu Glu Trp
1 5 10 15
Leu Gln Gly Arg Ala Cys Arg Pro Cys Gly Leu Leu Ala Ser Asp Ala
20 25 30
Ala Ala Leu Trp Phe Arg Gly Gly Ile Ser Ala Trp Glu Asp Ser Cys
35 40 45
Ala Val Ser Asn Ile Arg His Glu Ala Tyr Asn Cys His Leu Ser Val
50 55 60
Phe Leu Asn Arg Cys Ala Asn Glu Leu Thr Val Gln Phe Leu Ile Ile
65 70 75 80
Leu Ala Phe Gln Ile Met Leu Ser Cys Ala Val Ile Ala Pro Ala Val
85 90 95
Pro Val Phe Gln Arg Leu Thr Leu Lys Arg Ser Gly Arg Thr Ser Leu
100 105 110
Gly Ser Thr Gly Arg Leu His Phe Cys Lys *
115 120 122

<210> 1638
<211> 69
<212> PRT
<213> Homo sapiens

<400> 1638
Met Lys Arg Leu Arg Phe Val Leu Arg Val Phe Gln Met Thr Ala Phe
1 5 10 15
Ile Thr Gly Ala His Thr Ile Thr Asn Tyr Ser Asp Arg Arg Leu Tyr
20 25 30
Ile Ser Pro Leu Ser His Phe Phe Met Asn Ser Gly Ser Ser Ala Gln
35 40 45
Ser Val Leu Ser His Ser Tyr Val Ser Gln Ile Phe Phe Lys Asn Val
50 55 60
Ser Lys Tyr Phe *
65 68

<210> 1639

<211> 92
<212> PRT
<213> Homo sapiens

<400> 1639
Met Tyr Val Ala Gly Tyr Leu Val Ala Asn Ser Ala Ile Cys Gln Leu
1 5 10 15
Thr Gln His Ser Leu Val Lys Leu Leu Leu Gln Gly Cys Phe Leu Ile
20 25 30
Gly Ser Leu His Leu Cys Ile Cys Val Pro Met Cys Val Cys Val Cys
35 40 45
Glu Tyr Arg Ile Leu His Asp Ser Lys Ile Ser Phe Lys Tyr Leu Arg
50 55 60
Phe Thr Ile Leu Lys Arg Glu Asn Lys Asn Lys Val Leu Gln Lys Leu
65 70 75 80
Lys Lys Asn Leu Lys Ser Val His Thr Leu Ser *
85 90 91

<210> 1640
<211> 58
<212> PRT
<213> Homo sapiens

<400> 1640
Met Thr Ala Trp Phe Cys Ser Phe Leu Ser Ser His Trp Val Ile Lys
1 5 10 15
Leu Pro Arg Phe Leu Leu Leu Val Leu Pro Phe Phe Trp Gly Lys Lys
20 25 30
Phe Ser Leu Gly Leu Ile Ser Gln Phe Phe Ser Lys Ala Tyr Phe Tyr
35 40 45
Ser Ser Tyr His Asn Tyr Ile His Thr *
50 55 57

<210> 1641
<211> 459
<212> PRT
<213> Homo sapiens

<400> 1641
Met Ser Asp Leu Leu Ser Val Phe Leu His Leu Leu Leu Leu Phe Lys
1 5 10 15
Leu Val Ala Pro Val Thr Phe Arg His His Arg Tyr Asp Asp Leu Val
20 25 30
Arg Thr Leu Tyr Lys Val Gln Asn Glu Cys Pro Gly Ile Thr Arg Val
35 40 45
Tyr Ser Ile Gly Arg Ser Val Glu Gly Arg His Leu Tyr Val Leu Glu
50 55 60
Phe Ser Asp His Pro Gly Ile His Glu Pro Leu Glu Pro Glu Val Lys
65 70 75 80
Tyr Val Gly Asn Met His Gly Asn Glu Ala Leu Gly Arg Glu Leu Met
85 90 95
Leu Gln Leu Ser Glu Phe Leu Cys Glu Glu Phe Arg Asn Arg Asn Gln

```
            100                      105                     110
Arg Ile Val Gln Leu Ile Gln Asp Thr Arg Ile His Ile Leu Pro Ser
        115                     120                     125
Met Asn Pro Asp Gly Tyr Glu Val Ala Ala Ala Gln Gly Pro Asn Lys
    130                     135                     140
Pro Gly Tyr Leu Val Gly Arg Asn Asn Ala Asn Gly Val Asp Leu Asn
145                     150                     155                     160
Arg Asn Phe Pro Asp Leu Asn Thr Tyr Ile Tyr Tyr Asn Glu Lys Tyr
                165                     170                     175
Gly Gly Pro Asn His His Leu Pro Leu Pro Asp Asn Trp Lys Ser Gln
            180                     185                     190
Val Glu Pro Glu Thr Arg Ala Val Ile Arg Trp Met His Ser Phe Asn
        195                     200                     205
Phe Val Leu Ser Ala Asn Leu His Gly Gly Ala Val Val Ala Asn Tyr
    210                     215                     220
Pro Tyr Asp Lys Ser Phe Glu His Arg Val Arg Gly Val Arg Arg Thr
225                     230                     235                     240
Ala Ser Thr Pro Thr Pro Asp Asp Lys Leu Phe Gln Lys Leu Ala Lys
                245                     250                     255
Val Tyr Ser Tyr Ala His Gly Trp Met Phe Gln Gly Trp Asn Cys Gly
            260                     265                     270
Asp Tyr Phe Pro Asp Gly Ile Thr Asn Gly Ala Ser Trp Tyr Ser Leu
        275                     280                     285
Ser Lys Gly Met Gln Asp Phe Asn Tyr Leu His Thr Asn Cys Phe Glu
    290                     295                     300
Ile Thr Leu Glu Leu Ser Cys Asp Lys Phe Pro Pro Glu Glu Glu Leu
305                     310                     315                     320
Gln Arg Glu Trp Leu Gly Asn Arg Glu Ala Leu Ile Gln Phe Leu Glu
                325                     330                     335
Gln Val His Gln Gly Ile Lys Gly Met Val Leu Asp Glu Asn Tyr Asn
            340                     345                     350
Asn Leu Ala Asn Ala Val Ile Ser Val Ser Gly Ile Asn His Asp Val
        355                     360                     365
Thr Ser Gly Asp His Gly Asp Tyr Phe Arg Leu Leu Leu Pro Gly Ile
    370                     375                     380
Tyr Thr Val Ser Ala Thr Ala Pro Gly Tyr Asp Pro Glu Thr Val Thr
385                     390                     395                     400
Val Thr Val Gly Pro Ala Glu Pro Thr Leu Val Asn Phe His Leu Lys
                405                     410                     415
Arg Ser Ile Pro Gln Val Ser Pro Val Arg Arg Ala Pro Ser Arg Arg
            420                     425                     430
His Gly Val Arg Ala Lys Val Gln Pro Gln Pro Arg Lys Lys Glu Met
        435                     440                     445
Glu Met Arg Gln Leu Gln Arg Gly Pro Ala  *
    450                     455         458


     &lt;210&gt; 1642
     &lt;211&gt; 144
     &lt;212&gt; PRT
     &lt;213&gt; Homo sapiens


     &lt;400&gt; 1642
Met Ala Arg Cys Thr Leu Thr Leu Leu Lys Thr Met Leu Thr Glu Leu
  1               5                  10                  15
Leu Arg Gly Gly Ser Phe Glu Phe Lys Asp Met Arg Val Pro Ser Ala
            20                  25                  30
```

```
Leu Val Thr Leu His Met Leu Leu Cys Ser Ile Pro Leu Ser Gly Arg
        35                  40                  45
Leu Asp Ser Asp Glu Gln Lys Ile Gln Asn Asp Ile Ile Asp Ile Leu
    50                  55                  60
Leu Thr Phe Thr Gln Gly Val Asn Glu Lys Leu Thr Ile Ser Glu Glu
65                  70                  75                  80
Thr Leu Ala Asn Asn Thr Trp Ser Leu Met Leu Lys Glu Val Leu Ser
                85                  90                  95
Ser Ile Leu Lys Val Pro Glu Gly Phe Phe Ser Gly Leu Ile Leu Leu
            100                 105                 110
Ser Glu Leu Leu Pro Leu Pro Leu Pro Met Gln Thr Thr Gln Val Ser
        115                 120                 125
Leu Pro Tyr Asn Met His Leu Ile Asn Asp Cys Ser Asn Thr Phe  *
    130                 135                 140         143
```

```
<210> 1643
<211> 70
<212> PRT
<213> Homo sapiens
```

```
<400> 1643
Met Gly Arg Arg Trp Leu Phe Leu Ile Ala Cys Leu Arg Ser Ala Ser
 1               5                  10                  15
Ile Leu Ala Trp Ala Thr Trp Arg Asn Pro Val Ser Thr Lys Asn Lys
            20                  25                  30
Lys Leu Ala Ser His Asp Gly Pro His Leu Ala Val Pro Ala Ile Arg
        35                  40                  45
Glu Ala Glu Ala Gly Arg Trp Leu Lys Pro Arg Arg Arg Arg Leu Gln
    50                  55                  60
Arg Pro Lys Ile Ala Arg
65                  70
```

```
<210> 1644
<211> 82
<212> PRT
<213> Homo sapiens
```

```
<400> 1644
Met Gly Met Gly Thr Leu Ile Ile Met Asn Val Trp Val Leu Phe Ile
 1               5                  10                  15
Pro Thr Arg Leu Arg Ile Asp Gln Gln Pro Val His Ile Lys Pro Ser
            20                  25                  30
Met Arg Val Leu Asp Lys Trp Val Ser Ala Phe Val His Lys Gly Phe
        35                  40                  45
Thr Trp Gly Thr Ser Glu Arg Ile Asn Thr Gly Ser Ser Ser Asp Ile
    50                  55                  60
Thr Leu Gly Ile Leu Asn Lys Cys Gly Trp Ala Val Phe Cys Ala Ala
65                  70                  75                  80
Pro  *
81
```

<210> 1645
<211> 256
<212> PRT
<213> Homo sapiens

<400> 1645
Met Ala Ala Leu Thr Val Thr Leu Met Val Leu Ser Ser Pro Leu Ala
1 5 10 15
Leu Ala Gly Asp Thr Gln Pro Arg Phe Leu Trp Gln Gly Lys Tyr Lys
20 25 30
Cys His Phe Phe Asn Gly Thr Glu Arg Val Gln Phe Leu Glu Arg Leu
35 40 45
Phe Tyr Asn Gln Glu Glu Phe Val Arg Phe Asp Ser Asp Val Gly Glu
50 55 60
Tyr Arg Ala Val Thr Glu Leu Gly Arg Pro Val Ala Glu Ser Trp Asn
65 70 75 80
Ser Gln Lys Asp Ile Leu Glu Asp Arg Arg Gly Gln Val Asp Thr Val
85 90 95
Cys Arg His Asn Tyr Gly Val Gly Glu Ser Phe Thr Val Gln Arg Arg
100 105 110
Val His Pro Glu Val Thr Val Tyr Pro Ala Lys Thr Gln Pro Leu Gln
115 120 125
His His Asn Leu Leu Val Cys Ser Val Ser Gly Phe Tyr Pro Gly Ser
130 135 140
Ile Glu Val Arg Trp Phe Arg Asn Gly Gln Glu Glu Lys Ala Gly Val
145 150 155 160
Val Ser Thr Gly Leu Ile Gln Asn Gly Asp Trp Thr Phe Gln Thr Leu
165 170 175
Val Met Leu Glu Thr Val Pro Arg Ser Gly Glu Val Tyr Thr Cys Gln
180 185 190
Val Glu His Pro Ser Val Met Ser Pro Leu Thr Val Glu Trp Arg Ala
195 200 205
Arg Ser Glu Ser Ala Gln Ser Lys Met Leu Ser Gly Val Gly Gly Phe
210 215 220
Val Leu Gly Leu Leu Phe Leu Gly Ala Gly Leu Phe Ile Tyr Phe Arg
225 230 235 240
Asn Gln Lys Gly His Ser Gly Leu Gln Pro Thr Gly Phe Leu Ser *
245 250 255

<210> 1646
<211> 263
<212> PRT
<213> Homo sapiens

<400> 1646
Met Val Ala Trp Arg Ser Ala Phe Leu Val Cys Leu Ala Phe Ser Leu
1 5 10 15
Ala Thr Leu Val Gln Arg Gly Ser Gly Asp Phe Asp Asp Phe Asn Leu
20 25 30
Glu Asp Ala Val Lys Glu Thr Ser Ser Val Lys Gln Pro Trp Asp His
35 40 45
Thr Thr Thr Thr Thr Thr Asn Arg Pro Gly Thr Thr Arg Ala Pro Ala
50 55 60
Lys Pro Pro Gly Ser Gly Leu Asp Leu Ala Asp Ala Leu Asp Asp Gln
65 70 75 80

Asp Asp Gly Arg Arg Lys Pro Gly Ile Gly Gly Arg Glu Arg Trp Asn
85 90 95
His Val Thr Thr Thr Thr Lys Arg Pro Val Thr Thr Arg Ala Pro Ala
100 105 110
Asn Thr Leu Gly Asn Asp Phe Asp Leu Ala Asp Ala Leu Asp Asp Arg
115 120 125
Asn Asp Arg Asp Asp Gly Arg Arg Lys Pro Ile Ala Gly Gly Gly Gly
130 135 140
Phe Ser Asp Lys Asp Leu Glu Asp Ile Val Gly Gly Gly Glu Tyr Lys
145 150 155 160
Pro Asp Lys Gly Lys Gly Asp Gly Arg Tyr Gly Ser Asn Asp Asp Pro
165 170 175
Gly Ser Gly Met Val Ala Glu Pro Gly Thr Ile Ala Gly Val Ala Ser
180 185 190
Ala Leu Ala Met Ala Leu Ile Gly Ala Val Ser Ser Tyr Ile Ser Tyr
195 200 205
Gln Gln Lys Lys Phe Cys Phe Ser Ile Gln Gln Gly Leu Asn Ala Asp
210 215 220
Tyr Val Lys Gly Glu Asn Leu Glu Ala Val Val Cys Glu Glu Pro Gln
225 230 235 240
Val Lys Tyr Ser Thr Leu His Thr Gln Ser Ala Glu Pro Pro Pro Pro
245 250 255
Pro Glu Pro Ala Arg Ile *
260 262

<210> 1647
<211> 74
<212> PRT
<213> Homo sapiens

<400> 1647
Met Tyr Leu Leu Cys Trp Leu Tyr Ile Met Gly Val Leu Gly Ala Ser
1 5 10 15
Cys Asn Trp His Val Gly Val Pro Phe Pro Gly Thr His Trp Pro Arg
20 25 30
Ser Gln Asn His Leu Leu Trp Val Tyr Asn His Leu Asn Glu Leu Pro
35 40 45
Val Pro Ala Gly Arg Ser Ser Glu Gln Leu Tyr Leu Gly Tyr Thr Glu
50 55 60
Lys Tyr Gly Arg Arg Glu Arg Lys Ala *
65 70 73

<210> 1648
<211> 58
<212> PRT
<213> Homo sapiens

<400> 1648
Met Gly Leu Cys Gly Met Trp Val Leu Thr Ala Phe Leu Cys Glu Pro
1 5 10 15
Met Gly Phe Arg His Arg Val Cys Pro His Arg Cys Val Arg Gly Ser
20 25 30
Gly Arg Gly Ser Gly Cys Glu Cys Val Thr Met Trp Pro Cys Gly Ile

```
        35                  40                  45
Asn Ala Met Thr Gly Gly Phe Trp Val  *
     50                  55      57


     <210> 1649
     <211> 90
     <212> PRT
     <213> Homo sapiens


     <400> 1649
Met Gly Val Leu Leu Val Ser Met Val Val Leu Phe Ile Phe Ala Ile
 1               5                  10                  15
Leu Cys Ile Phe Ile Arg Asn Arg Ile Leu Glu Ile Val Tyr Ala Ser
            20                  25                  30
Leu Gly Ala Leu Leu Phe Thr Cys Phe Leu Ala Val Asp Thr Gln Leu
        35                  40                  45
Leu Leu Gly Asn Lys Gln Leu Ser Leu Ser Pro Glu Glu Tyr Val Phe
    50                  55                  60
Ala Ala Leu Asn Leu Tyr Thr Asp Ile Ile Asn Ile Phe Leu Tyr Ile
65                  70                  75                  80
Leu Thr Ile Ile Gly Arg Ala Lys Glu  *
                85              89


     <210> 1650
     <211> 113
     <212> PRT
     <213> Homo sapiens


     <400> 1650
Met Ala Leu Gly Val Pro Ile Ser Val Tyr Leu Leu Phe Asn Ala Met
 1               5                  10                  15
Thr Ala Leu Thr Glu Glu Ala Ala Val Thr Val Thr Pro Pro Ile Thr
            20                  25                  30
Ala Gln Gln Gly Asn Trp Thr Val Asn Lys Thr Glu Ala Asp Asn Ile
        35                  40                  45
Glu Gly Pro Ile Ala Leu Lys Phe Ser His Leu Cys Leu Glu Asp His
    50                  55                  60
Asn Ser Tyr Cys Ile Asn Gly Ala Cys Ala Phe His His Glu Leu Glu
65                  70                  75                  80
Lys Ala Ile Cys Arg Cys Phe Thr Gly Tyr Thr Gly Glu Arg Cys Leu
                85                  90                  95
Lys Leu Lys Ser Pro Tyr Asn Val Cys Ser Gly Glu Arg Arg Pro Leu
            100                 105                 110     112
 *


     <210> 1651
     <211> 50
     <212> PRT
     <213> Homo sapiens
```

<400> 1651
Met Phe Ile Lys Phe Leu Arg Ile Leu Ile Ser Leu Gln Cys Ser Ser
1 5 10 15
Phe Lys Phe Thr Val Thr Ala Lys Val Leu Phe Met Thr Tyr Lys Lys
20 25 30
Arg Ala Gln Ser Asp Phe Phe Leu Val Phe Val Asp Arg Glu Arg Ser
35 40 45
Pro *
49

<210> 1652
<211> 121
<212> PRT
<213> Homo sapiens

<400> 1652
Met Ser Arg Ala Gly Met Leu Gly Val Val Cys Ala Leu Leu Val Trp
1 5 10 15
Ala Tyr Leu Ala Val Gly Lys Leu Val Val Arg Met Thr Phe Thr Glu
20 25 30
Leu Cys Thr His His Pro Trp Ser Leu Arg Cys Glu Ser Phe Cys Arg
35 40 45
Ser Arg Val Thr Ala Cys Leu Pro Ala Pro Ala Pro Trp Leu Arg Pro
50 55 60
Phe Leu Cys Pro Met Leu Phe Ser Asp Arg Asn Pro Val Glu Cys His
65 70 75 80
Leu Phe Gly Glu Ala Val Ser Asp Pro Val Cys Lys Gly Leu Leu Pro
85 90 95
His Tyr Phe Trp His Pro Thr Phe Phe Pro Val Lys Ala Asn Cys Leu
100 105 110
Val Ser Phe Cys Pro Thr Thr Val *
115 120

<210> 1653
<211> 111
<212> PRT
<213> Homo sapiens

<400> 1653
Met Trp Ser Leu Trp Ile Trp Val Asp Gln His Gln Ala Arg Leu Ile
1 5 10 15
Pro Ser Pro Gln Val Leu Leu Leu Leu Leu Arg Glu Thr Pro Ser Thr
20 25 30
Ala Ala Ala Val Ala Gly Trp Leu Val Val Ala Ser Met Ala Leu Leu
35 40 45
Gln Leu His Ala Val Gly Gly Val Ala Leu Thr Ser Ser His Pro Phe
50 55 60
Met Trp Ala Thr Gly Glu Glu Leu Arg Lys Pro Pro Trp Gln Gly Ser
65 70 75 80
Ala Gly Ser Ala Ser Gly Val Glu Glu Leu Thr Gly Lys His Ser Cys
85 90 95
Pro Gly Pro Glu Glu Pro Ala Thr Val Gln Lys Ala Pro Ala *

100 105 110

<210> 1654
<211> 150
<212> PRT
<213> Homo sapiens

<400> 1654
Met Trp Ile Cys Arg Val Lys Gln Ala Trp Leu Pro Pro Leu Leu Ser
1 5 10 15
Pro Leu Gly Pro Pro Thr Pro Trp Asp Pro Phe Tyr Ala Ala Pro Ser
20 25 30
Pro Pro Val Trp Val Gly Ser Gly Tyr Trp Tyr Arg Gly Leu Leu Ser
35 40 45
Pro Pro Asp Gly Gly Gln Gly Ser Phe Pro Pro His Leu Cys Pro Gln
50 55 60
Cys Pro Val Gln Ala Gln Ala Gln Ile Gly Pro Tyr Phe Arg Glu Leu
65 70 75 80
Gly Glu Pro Pro Ser Glu Thr Lys Trp Tyr Leu Asn Ser His Ser His
85 90 95
His Arg Ala Ala Gly Thr Gln Arg Arg Leu Arg Cys Leu Gln His Leu
100 105 110
Leu Gly Gly Gly Gly Pro Gly Ile Gly Ser Glu Ser Pro Asn Glu Gly
115 120 125
Pro Gly Gln Val Thr His Ala Cys Asn Leu Ser Thr Leu Gly Gly Lys
130 135 140
Asp Val Arg Ile Thr *
145 149

<210> 1655
<211> 68
<212> PRT
<213> Homo sapiens

<400> 1655
Met Ser Arg Asn Leu Arg Thr Ala Leu Ile Phe Gly Gly Phe Ile Ser
1 5 10 15
Leu Ile Gly Ala Ala Phe Tyr Pro Ile Tyr Phe Arg Pro Leu Met Arg
20 25 30
Leu Glu Glu Tyr Lys Lys Glu Gln Ala Ile Asn Arg Ala Gly Ile Val
35 40 45
Gln Glu Asp Val Gln Pro Pro Gly Leu Lys Val Trp Ser Asp Pro Phe
50 55 60
Gly Arg Lys *
65 67

<210> 1656
<211> 61
<212> PRT
<213> Homo sapiens

<400> 1656

```
Met His Lys Tyr Leu Cys Val Phe Glu Tyr Leu Ser Asn Leu Ser Lys
 1               5                  10                  15
Cys Met Arg Leu Tyr Leu Ile Leu Leu Ala Ser Ile Cys Met Tyr Leu
            20                  25                  30
Cys Val Ala Arg Arg Val Phe Leu Phe Ala Ser Val Ser Thr Gln Gly
        35                  40                  45
Lys Ser Leu Met Tyr Ser Thr Gln Lys Val Val Lys  *
    50                  55                  60
```

<210> 1657
<211> 80
<212> PRT
<213> Homo sapiens

<400> 1657

```
Met Asn Trp Gln His Ser Thr Met Tyr Leu Phe Phe Ala Val Ser Gly
 1               5                  10                  15
Ile Val Asp Met Leu Thr Tyr Leu Val Ser His Val Pro Leu Gly Val
            20                  25                  30
Asp Arg Leu Val Met Gly Cys Gly Lys Tyr Ser Trp Lys Val Ser Ser
        35                  40                  45
Ser Thr Thr Thr Ser Thr Thr Gly Leu Arg Trp Thr Ser Thr Ser Thr
    50                  55                  60
His Ser Cys Cys Met Leu Cys Ser Glu Gly Val Leu Val Ser Pro  *
65                  70                  75                  79
```

<210> 1658
<211> 160
<212> PRT
<213> Homo sapiens

<400> 1658

```
Met Ala Phe Leu Leu Tyr His Leu Val Tyr His Ile Pro Pro Met Ala
 1               5                  10                  15
Pro Val Ser Phe Val Phe Glu Thr Lys Ser Arg Ser Ala Ala Gln Ala
            20                  25                  30
Gly Val Gln Trp His Asp Pro Gly Ser Pro Gln Pro Leu Pro Pro Arg
        35                  40                  45
Phe Lys Arg Phe Ser Cys His Gly Leu Asn Ile Lys Phe Ala Phe Phe
    50                  55                  60
Ser His Leu Lys Glu Leu His Leu Asp Ser Gly His Cys Phe Ile Phe
65                  70                  75                  80
Ile Arg Leu Val Lys Gly Ala Val Cys Leu Ile His Val Gln Ile Arg
                85                  90                  95
Ile Pro Ser Ala Asp Glu Asp Ile Thr Ile Leu Phe Phe Ile Val Ser
            100                 105                 110
Lys His Phe Leu Glu Ser Val Phe Lys Met Leu Gln Trp Ser Gln Met
        115                 120                 125
Thr Leu Ala Thr Val Lys Thr Thr Phe Ile Gly Leu Asn Glu Phe Ile
    130                 135                 140
Cys Ser Pro Ser Thr Leu Pro Ser Gly Lys Lys Asn Gly Leu Ile  *
```

145 150 155 159

<210> 1659
<211> 90
<212> PRT
<213> Homo sapiens

<400> 1659
Met Trp Arg Leu Pro His Ser Gln Phe Ile His Ile Val Ile Leu Pro
1 5 10 15
Leu Lys Val Phe Leu Phe Leu Phe Cys Phe Leu Arg Trp Ser Phe Ser
20 25 30
Leu Val Ala Gln Ala Gly Val Gln Trp Arg Asp Leu Gly Pro Leu Gln
35 40 45
Pro Pro Pro Pro Arg Leu Lys Arg Phe Phe Cys Leu Ser Leu Pro Ser
50 55 60
Ser Trp Asp Tyr Arg His Ser Pro Pro His Pro Ala Asn Phe Tyr Thr
65 70 75 80
Phe Gly Arg Asp Gly Val Ser Pro Cys *
85 89

<210> 1660
<211> 56
<212> PRT
<213> Homo sapiens

<400> 1660
Met Cys Ala His Leu Val Cys Val Lys Trp Cys Leu Val Ile Leu Ile
1 5 10 15
Cys Ile Phe Gln Asn Thr Asn Glu Val Glu Gln Leu Ile Leu Cys Val
20 25 30
Leu Leu Ile Pro Leu Ser Ser Ser Met Thr Asp Leu Phe Leu Ser Leu
35 40 45
Cys Val Cys Val Phe Cys Tyr *
50 55

<210> 1661
<211> 74
<212> PRT
<213> Homo sapiens

<400> 1661
Met Leu Gly Met Ile Ser Met Leu Leu Asn Ala Leu Lys Leu Leu Val
1 5 10 15
Tyr Leu Thr Glu Cys Cys Met Ala Leu Glu Glu Arg Val His Ser Val
20 25 30
Leu Ile Gly Trp Ser Val Ser Phe Lys Arg Ile Gln Arg Gln Leu Asn
35 40 45
Gln Val Gly Leu Ile Glu Phe Phe Lys Met Val Leu Cys Ser Asn Thr
50 55 60

Asp Gly Thr Glu Gly His Tyr Pro Lys *
65 70 73

<210> 1662
<211> 271
<212> PRT
<213> Homo sapiens

<400> 1662
Met Gly Leu Gly Gln Pro Gln Ala Trp Leu Leu Gly Leu Pro Thr Ala
1 5 10 15
Val Val Tyr Gly Ser Leu Ala Leu Phe Thr Thr Ile Leu His Asn Val
20 25 30
Phe Leu Leu Tyr Tyr Val Asp Thr Phe Val Ser Val Tyr Lys Ile Asn
35 40 45
Lys Met Ala Phe Trp Val Gly Glu Thr Val Phe Leu Leu Trp Asn Ser
50 55 60
Leu Asn Asp Pro Leu Phe Gly Trp Leu Ser Asp Arg Gln Phe Leu Ser
65 70 75 80
Ser Gln Pro Arg Ser Gly Ala Gly Leu Ser Ser Arg Ala Val Val Leu
85 90 95
Ala Arg Val Gln Ala Leu Gly Trp His Gly Pro Leu Leu Ala Leu Ser
100 105 110
Phe Leu Ala Phe Trp Val Pro Trp Ala Pro Ala Gly Leu Gln Phe Leu
115 120 125
Leu Cys Leu Cys Leu Tyr Asp Gly Phe Leu Thr Leu Val Asp Leu His
130 135 140
His His Ala Leu Leu Ala Asp Leu Ala Leu Ser Ala His Asp Arg Thr
145 150 155 160
His Leu Asn Phe Tyr Cys Ser Leu Phe Ser Ala Ala Gly Ser Leu Ser
165 170 175
Val Phe Ala Ser Tyr Ala Phe Trp Asn Lys Glu Asp Phe Ser Ser Phe
180 185 190
Arg Ala Phe Cys Val Thr Leu Ala Val Ser Ser Gly Leu Gly Phe Leu
195 200 205
Gly Ala Thr Gln Leu Leu Arg Arg Arg Val Glu Ala Ala Arg Lys Asp
210 215 220
Pro Gly Cys Ser Gly Leu Val Val Asp Ser Gly Leu Cys Gly Glu Glu
225 230 235 240
Leu Leu Val Gly Ser Glu Glu Ala Asp Ser Ile Thr Leu Gly Arg Tyr
245 250 255
Leu Arg Gln Leu Ala Arg His Arg Asn Phe Leu Cys Phe Ser *
260 265 270

<210> 1663
<211> 53
<212> PRT
<213> Homo sapiens

<400> 1663
Met Pro His Ile Gln Thr Leu Leu Arg Thr Leu Phe Ala Ser His Leu
1 5 10 15
Leu Val Ser Leu Trp Gln Ser Glu Pro Met Ala Lys Pro Arg Met Arg

```
            20                  25                  30
Lys Tyr Asn Thr Ser Ser Glu Tyr Leu Ser Glu Leu Asp Thr Glu Ala
        35                  40                  45
Ser Arg Val Ser  *
    50      52
```

<210> 1664
<211> 271
<212> PRT
<213> Homo sapiens

<400> 1664

```
Met Gly Leu Gly Gln Pro Gln Ala Trp Leu Leu Gly Leu Pro Thr Ala
 1               5                  10                  15
Val Val Tyr Gly Ser Leu Ala Leu Phe Thr Thr Ile Leu His Asn Val
            20                  25                  30
Phe Leu Leu Tyr Tyr Val Asp Thr Phe Val Ser Val Tyr Lys Ile Asn
        35                  40                  45
Lys Met Ala Phe Trp Val Gly Glu Thr Val Phe Leu Leu Trp Asn Ser
    50                  55                  60
Leu Asn Asp Pro Leu Phe Gly Trp Leu Ser Asp Arg Gln Phe Leu Ser
65                  70                  75                  80
Ser Gln Pro Arg Ser Gly Ala Gly Leu Ser Ser Arg Ala Val Val Leu
                85                  90                  95
Ala Arg Val Gln Ala Leu Gly Trp His Gly Pro Leu Leu Ala Leu Ser
            100                 105                 110
Phe Leu Ala Phe Trp Val Pro Trp Ala Pro Ala Gly Leu Gln Phe Leu
        115                 120                 125
Leu Cys Leu Cys Leu Tyr Asp Gly Phe Leu Thr Leu Val Asp Leu His
    130                 135                 140
His His Ala Leu Leu Ala Asp Leu Ala Leu Ser Ala His Asp Arg Thr
145                 150                 155                 160
His Leu Asn Phe Tyr Cys Ser Leu Phe Ser Ala Ala Gly Ser Leu Ser
                165                 170                 175
Val Phe Ala Ser Tyr Ala Phe Trp Asn Lys Glu Asp Phe Ser Ser Phe
            180                 185                 190
Arg Ala Phe Cys Val Thr Leu Ala Val Ser Ser Gly Leu Gly Phe Leu
        195                 200                 205
Gly Ala Thr Gln Leu Leu Arg Arg Arg Val Glu Ala Ala Arg Lys Asp
    210                 215                 220
Pro Gly Cys Ser Gly Leu Val Val Asp Ser Gly Leu Cys Gly Glu Glu
225                 230                 235                 240
Leu Leu Val Gly Ser Glu Glu Ala Asp Ser Ile Thr Leu Gly Arg Tyr
                245                 250                 255
Leu Arg Gln Leu Ala Arg His Arg Asn Phe Leu Cys Phe Ser  *
            260                 265                 270
```

<210> 1665
<211> 284
<212> PRT
<213> Homo sapiens

<400> 1665

```
Met Asp Glu Lys Ser Asn Lys Leu Leu Leu Ala Leu Val Met Leu Phe
 1               5                  10                  15
Leu Phe Ala Val Ile Val Leu Gln Tyr Val Cys Pro Gly Thr Glu Cys
            20                  25                  30
Gln Leu Leu Arg Leu Gln Ala Phe Ser Ser Pro Val Pro Asp Pro Tyr
        35                  40                  45
Arg Ser Glu Asp Glu Ser Ser Ala Arg Phe Val Pro Arg Tyr Asn Phe
    50                  55                  60
Thr Arg Gly Asp Leu Leu Arg Lys Val Asp Phe Asp Ile Lys Gly Asp
65                  70                  75                  80
Asp Leu Ile Val Phe Leu His Ile Gln Lys Thr Gly Gly Thr Thr Phe
                85                  90                  95
Gly Arg His Leu Val Arg Asn Ile Gln Leu Glu Gln Pro Cys Glu Cys
            100                 105                 110
Arg Val Gly Gln Lys Lys Cys Thr Cys His Arg Pro Gly Lys Arg Glu
        115                 120                 125
Thr Trp Leu Phe Ser Arg Phe Ser Thr Gly Trp Ser Cys Gly Leu His
    130                 135                 140
Ala Asp Trp Thr Glu Leu Thr Ser Cys Val Pro Ser Val Gly Asp Gly
145                 150                 155                 160
Lys Arg Asp Ala Arg Leu Arg Pro Ser Arg Trp Arg Ile Phe His Ile
                165                 170                 175
Leu Tyr Ala Ala Cys Thr Asp Ile Arg Gly Ser Pro Asn Thr Asn Ala
            180                 185                 190
Gly Ala Asn Ser Pro Ser Phe Thr Lys Thr Arg Asn Thr Ser Lys Ser
        195                 200                 205
Trp Lys Asn Phe His Tyr Ile Thr Ile Leu Gln Asp Pro Gly Ala Arg
    210                 215                 220
Ser Leu Ser Glu Trp Arg Pro Val Leu Lys Arg Gly Thr Leu Glu Gly
225                 230                 235                 240
Leu Leu Ala Cys Trp Pro Trp Lys Ala Pro Pro Pro Leu Lys Lys Leu
                245                 250                 255
Ser Thr Trp Tyr Pro Gly Glu Glu Leu Val Trp Leu Ala Pro Leu Gln
            260                 265                 270
Lys Ile Ile Gly Leu Ala Leu Leu Ile Tyr Pro  *
        275                 280         283
```

```
<210> 1666
<211> 67
<212> PRT
<213> Homo sapiens


<400> 1666
Met Thr Leu Val Leu Phe Leu Val Leu Ala Leu Met Ile Thr Ile Cys
 1               5                  10                  15
Ile Leu Ser Tyr His Ser His Leu Leu Ile Asn Ser Asn Leu Ile Pro
            20                  25                  30
Val Lys Tyr Arg Asn Phe Pro Ser Ile Leu Leu His Phe Leu His Leu
        35                  40                  45
Trp Leu Ser Phe Cys His Ile Ser His Met His Ile Cys His Asn Leu
    50                  55                  60
Leu Ile  *
 65  66
```

<210> 1667
<211> 79
<212> PRT
<213> Homo sapiens

<400> 1667
Met Asn Thr His Trp Asn Ile Leu Pro Val Glu Arg Ser Cys Pro Leu
1 5 10 15
Trp Ile Ser Ser Glu Leu Ser Tyr Cys Ser Ile Lys Leu Leu Phe Ile
20 25 30
Leu Leu Thr Leu His Leu Pro Ala Tyr Leu Ile Leu Pro Gly His Lys
35 40 45
Ile Arg Thr Gln Asp Leu Pro Asn Glu Ala Asn Arg Ala Val Thr Gln
50 55 60
Thr Gly Leu Arg His Ala Leu Tyr Gln Ser Ile Ser Cys Trp *
65 70 75 78

<210> 1668
<211> 54
<212> PRT
<213> Homo sapiens

<400> 1668
Met Trp Gly Leu Leu Ile Pro Cys Ile Leu Gly Cys Met Lys Leu Pro
1 5 10 15
His Asn Leu Leu Met Leu Phe Ser Leu Glu Thr Phe Leu Thr Leu Arg
20 25 30
Phe Ile Leu Asp Ser Phe Tyr Ser Tyr Val Phe Lys Pro Thr Asn Lys
35 40 45
Arg Phe Cys Asn Ile *
50 53

<210> 1669
<211> 119
<212> PRT
<213> Homo sapiens

<400> 1669
Met Met Ala Gly Ile Arg Ala Leu Phe Met Tyr Leu Trp Leu Gln Leu
1 5 10 15
Asp Trp Val Ser Arg Gly Glu Ser Val Gly Leu His Leu Pro Thr Leu
20 25 30
Ser Val Gln Glu Gly Asp Asn Ser Ile Ile Asn Cys Ala Tyr Ser Asn
35 40 45
Ser Ala Ser Asp Tyr Phe Ile Trp Tyr Lys Gln Glu Ser Gly Lys Gly
50 55 60
Pro Gln Phe Ile Ile Asp Ile Arg Ser Asn Met Asp Lys Arg Gln Gly
65 70 75 80
Gln Arg Val Thr Val Leu Leu Asn Lys Thr Val Lys His Leu Ser Leu
85 90 95
Gln Ile Ala Ala Thr Gln Pro Gly Asp Ser Ala Val Tyr Phe Cys Ala
100 105 110

```
Glu Ile Pro Glu Gln Arg  *
        115          118


     <210> 1670
     <211> 116
     <212> PRT
     <213> Homo sapiens


     <400> 1670
Met Cys Leu Leu Cys Cys Glu Cys Leu Phe His Leu Trp Lys Arg Ile
 1               5                  10                  15
Asn Trp Trp Gln Gly Phe Cys Ser Phe Tyr Leu Leu Leu Trp Val Gly
            20                  25                  30
Leu Leu Ser Phe Pro Pro Asp Pro Pro Trp Lys Ser Phe Thr Pro Ala
        35                  40                  45
Ile Leu Phe Leu Ala Trp Gly Thr Gly Ser Ser Pro Gly Arg His Arg
    50                  55                  60
Phe Ser Leu Pro Thr Asp Arg Arg Pro Ser Ala His Ser Pro Phe Leu
65                  70                  75                  80
Ser Thr Leu Gln His Ser Ile Arg Thr Leu Phe His Ser Pro Ile Arg
                85                  90                  95
Ser Ser Arg Phe Ala Phe Val Ser Ser Leu His Ser Tyr Thr Ser Ile
            100                 105                 110
Pro Ser Leu Pro
        115 116


     <210> 1671
     <211> 70
     <212> PRT
     <213> Homo sapiens


     <400> 1671
Met Ser His Cys Gly Leu Leu Phe Leu Val Val Thr Trp Leu Leu Ser
 1               5                  10                  15
Phe Ile Phe Leu Val Cys Lys Met Arg Ile Thr Phe Leu Phe Cys Leu
            20                  25                  30
Leu Thr Val Asp Met Lys Pro Asn Lys Val Leu Tyr Met Lys Cys Phe
        35                  40                  45
Lys Cys Ile Ile Leu Leu Ser Cys Tyr Pro Leu Lys Phe Leu Val Ile
    50                  55                  60
Arg Asn Phe Glu Ile  *
65              69


     <210> 1672
     <211> 263
     <212> PRT
     <213> Homo sapiens


     <400> 1672
Met Arg Val Leu Cys Ala Phe Pro Glu Ala Met Pro Ser Ser Asn Ser
```

```
 1               5                   10                  15
Arg Pro Pro Ala Cys Leu Ala Pro Gly Ala Leu Tyr Leu Ala Leu Leu
            20                  25                  30
Leu His Leu Ser Leu Ser Ser Gln Ala Gly Asp Arg Arg Pro Leu Pro
        35                  40                  45
Val Asp Arg Ala Ala Gly Leu Lys Glu Lys Thr Leu Ile Leu Leu Asp
    50                  55                  60
Val Ser Thr Lys Asn Pro Val Arg Thr Val Asn Glu Asn Phe Leu Ser
65                  70                  75                      80
Leu Gln Leu Asp Pro Ser Ile Ile His Asp Gly Trp Leu Asp Phe Leu
                85                  90                  95
Ser Ser Lys Arg Leu Val Thr Leu Ala Arg Gly Leu Ser Pro Ala Phe
            100                 105                 110
Leu Arg Phe Gly Gly Lys Arg Thr Asp Phe Leu Gln Phe Gln Asn Leu
        115                 120                 125
Arg Asn Pro Ala Lys Ser Arg Gly Gly Pro Gly Pro Asp Tyr Tyr Leu
    130                 135                 140
Lys Asn Tyr Glu Asp Asp Ile Val Arg Ser Asp Val Ala Leu Asp Lys
145                 150                 155                     160
Gln Lys Gly Cys Lys Ile Ala Gln His Pro Asp Gly Met Leu Glu Pro
                165                 170                 175
Pro Arg Glu Lys Ala Ala Gln Met His Leu Val Leu Leu Lys Glu Gln
            180                 185                 190
Phe Ser Asn Thr Tyr Ser Asn Leu Ile Leu Thr Glu Pro Asn Asn Tyr
        195                 200                 205
Arg Thr Met His Gly Arg Ala Val Asn Gly Ser Gln Leu Gly Lys Asp
    210                 215                 220
Tyr Ile Gln Leu Lys Ser Leu Leu Gln Pro Ile Arg Ile Tyr Ser Arg
225                 230                 235                     240
Ala Ser Leu Tyr Gly Pro Asn Ile Val Arg Pro Arg Lys Asn Val Ile
                245                 250                 255
Ala Leu Leu Asp Gly Leu  *
            260     262
```

<210> 1673
<211> 156
<212> PRT
<213> Homo sapiens

<400> 1673

```
Met Lys Trp Lys Thr Gly Val Ala Ile Phe Val Val Val Val Val Tyr
 1               5                   10                  15
Leu Val Thr Gly Gly Leu Val Phe Arg Ala Leu Glu Gln Pro Phe Glu
            20                  25                  30
Ser Ser Gln Lys Asn Thr Ile Ala Leu Glu Lys Ala Glu Phe Leu Arg
        35                  40                  45
Asp His Val Cys Val Ser Pro Gln Glu Leu Glu Thr Leu Ile Gln His
    50                  55                  60
Ala Leu Asp Ala Asp Asn Ala Gly Val Ser Pro Ile Gly Asn Ser Ser
65                  70                  75                      80
Asn Asn Ser Ser His Trp Asp Leu Gly Ser Ala Phe Phe Phe Ala Gly
                85                  90                  95
Thr Val Ile Thr Thr Ile Gly Tyr Gly Asn Ile Ala Pro Ser Thr Glu
            100                 105                 110
Gly Gly Lys Ile Phe Cys Ile Leu Tyr Ala Ile Phe Gly Phe Pro Leu
        115                 120                 125
```

```
Phe Gly Phe Leu Leu Ala Gly Ile Glu Asp Gln Leu Gly Thr Ile Phe
    130                 135                 140
Gly Lys Ser Ile Ala Arg Val Glu Lys Val Phe  *
145                 150                 155
```

<210> 1674
<211> 83
<212> PRT
<213> Homo sapiens

<400> 1674

```
Met Cys Cys Val Ile Cys Ser Lys Gln Tyr Val Leu Leu Ser Ile Leu
  1               5                  10                  15
Leu Cys Leu Leu Ala Ser Gly Ser Val Asp Phe Phe Leu Leu Pro His
             20                  25                  30
Ser Val Leu Ala Asp Asp Asp Gly Ile Lys Val Val Lys Val Thr Phe
         35                  40                  45
Asn Lys Gln Asp Ser Leu Val Ile Leu Thr Ile Met Val Ser Leu Thr
     50                  55                  60
Val Ser Phe Pro Gly Leu Cys Thr Cys Gln Ala Gly Thr Gln Asp Thr
 65                  70                  75                  80
Tyr Thr  *
     82
```

<210> 1675
<211> 54
<212> PRT
<213> Homo sapiens

<400> 1675

```
Met Val His Cys Leu Ile Cys Met Trp Thr Cys Trp Pro Thr Gly Ala
  1               5                  10                  15
Ile Leu His Arg Val Cys Arg Thr His Trp Pro Arg Gly Val Ser His
             20                  25                  30
Thr His Val Trp Met His Trp Pro Thr Cys Val Val Ser Arg Leu Phe
         35                  40                  45
Val Asp Val Leu Gly  *
     50          53
```

<210> 1676
<211> 119
<212> PRT
<213> Homo sapiens

<400> 1676

```
Met Gly Val Met Ala Met Leu Met Leu Pro Leu Leu Leu Leu Gly Ile
  1               5                  10                  15
Ser Gly Leu Leu Phe Ile Tyr Gln Glu Val Ser Arg Leu Trp Ser Lys
             20                  25                  30
Ser Ala Val Gln Asn Lys Val Val Val Ile Thr Asp Ala Ile Ser Gly
```

35 40 45
Leu Gly Lys Glu Cys Ala Arg Val Phe His Thr Gly Gly Ala Arg Leu
50 55 60
Val Leu Cys Gly Lys Asn Trp Glu Arg Leu Glu Asn Leu Tyr Asp Ala
65 70 75 80
Leu Ile Ser Val Ala Asp Pro Ser Lys Thr Phe Thr Pro Lys Leu Val
85 90 95
Leu Leu Asp Leu Ser Asp Ile Ser Cys Val Pro His Val Ala Lys Glu
100 105 110
Ala Leu Asp Cys Tyr Gly *
115 118

<210> 1677
<211> 49
<212> PRT
<213> Homo sapiens

<400> 1677
Met Arg Tyr Lys Cys Val Leu Ser Lys Ile Leu Trp Phe Cys Pro Trp
1 5 10 15
Lys Tyr Val Trp Lys Asn Ser Phe Phe Asn Leu Glu Gly Met Phe Met
20 25 30
Phe Ile Glu Val Thr Cys Arg His Tyr Ser Thr Cys Gly Ile Phe Lys
35 40 45 48
*

<210> 1678
<211> 127
<212> PRT
<213> Homo sapiens

<400> 1678
Met Gln Thr Lys Gly Gly Gln Thr Trp Ala Arg Arg Ala Leu Leu Leu
1 5 10 15
Gly Ile Leu Trp Ala Thr Ala His Leu Pro Leu Ser Gly Thr Ser Leu
20 25 30
Pro Gln Arg Leu Pro Arg Ala Thr Gly Asn Ser Thr Gln Cys Val Ile
35 40 45
Ser Pro Ser Ser Glu Phe Pro Glu Gly Phe Phe Thr Arg Gln Glu Arg
50 55 60
Arg Asp Gly Gly Ile Ile Ile Tyr Phe Leu Ile Ile Val Tyr Met Phe
65 70 75 80
Met Ala Ile Ser Ile Val Cys Asp Glu Tyr Phe Leu Pro Ser Leu Glu
85 90 95
Ile Ile Ser Glu Tyr Ile Gly Asn Lys Lys Glu Met Gln Val Leu Ile
100 105 110
Pro Gly Arg Ile Val Ser Lys Leu Lys Lys Leu Gly Phe Lys *
115 120 125 126

<210> 1679

<211> 49
<212> PRT
<213> Homo sapiens

<400> 1679
Met Ile Phe Phe Ile Lys Ala Pro Leu Tyr Leu Leu Gln Ser Met Met
1 5 10 15
Asp Cys Leu Tyr Ala Arg Arg Ile Pro Cys Ile Thr Asp Cys Ala Met
20 25 30
Ala Glu Ile Glu Lys Leu Gly Gln Lys Tyr Pro Val Ala Leu Arg Ile
35 40 45
Ala
49

<210> 1680
<211> 58
<212> PRT
<213> Homo sapiens

<400> 1680
Met Val Tyr Glu Val Phe Ile Asn Lys Ala Asn Ile Leu Leu Leu Leu
1 5 10 15
Phe Leu Arg Gln Ser Leu Ala Val Leu Pro Arg Leu Glu Cys Ser Gly
20 25 30
Ala Ile Ser Ala Arg Cys Asn Leu His Leu Arg Ile Pro Pro Asp Phe
35 40 45
His Arg Ser Thr Met Gly Gly Gly Gly Gly
50 55 58

<210> 1681
<211> 49
<212> PRT
<213> Homo sapiens

<400> 1681
Met Leu Ser Gly Trp Val Gln Cys Pro Leu Leu Gln Arg Val His Phe
1 5 10 15
Tyr Ala Phe Ser Val Gly Pro Phe His Arg Lys Ile Trp Gly Asp Val
20 25 30
Ser Phe Pro Leu Thr Phe Tyr Phe Lys Asn Leu Gln Thr Gln Lys Ser
35 40 45 48
*

<210> 1682
<211> 78
<212> PRT
<213> Homo sapiens

<400> 1682
Met Thr Gly Leu Phe Leu His His Asn Pro Gly Ile Leu Leu Ala Pro
1 5 10 15
Ser Val Leu Asp Leu Leu Phe Pro Gly Ser His Ile Phe Ile Phe Ser
20 25 30
Leu Phe Leu Ser Leu Cys Pro Cys Phe Gly Asp Thr Ile Leu Val Ala
35 40 45
Pro Ser Asp Lys Val Tyr Lys Asp Thr Phe Ile Ile Lys Ile Tyr Pro
50 55 60
Tyr Cys Ile Phe Glu Asn Phe Phe Thr Phe Leu Phe Thr *
65 70 75 77

<210> 1683
<211> 52
<212> PRT
<213> Homo sapiens

<400> 1683
Met Ser Leu Gly Ser Ile Asn His Phe Leu Phe Phe Ile Gln Leu Leu
1 5 10 15
Val Leu Lys Asn Ser Tyr Cys Met Leu Leu Lys Met Lys Gln Asn Lys
20 25 30
Lys Leu Lys Lys Ile Met Cys Leu Leu Phe Leu Met Leu Ser Ser Tyr
35 40 45
His Leu Ile *
50 51

<210> 1684
<211> 165
<212> PRT
<213> Homo sapiens

<400> 1684
Met Pro Ala Pro Pro Leu Pro Gly Gly Trp Asn Thr Trp Gly Pro Ser
1 5 10 15
Leu Ser Leu Pro Leu Leu Leu Leu Gly Ala Val Ala Met Ala Leu Gly
20 25 30
Val Arg Pro Pro Gly Gln Val Gly Leu Ser Pro Ile Ala Thr Ala Ser
35 40 45
Thr Val Gly Val Pro Arg Cys Leu Gln Thr Ala Phe Arg Gly Asp Ala
50 55 60
Gly Trp His Ser Cys Ala Gln Gln Gly Ala Cys Val Ala Leu His Pro
65 70 75 80
Ser Glu Arg Arg Leu Gly Ile Ser Asp Glu Ala His Ser Arg Ser Arg
85 90 95
Trp Gly Gly Glu Asp Ser Pro Ser Pro Leu Thr Gly Pro Pro Leu Ser
100 105 110
Pro Ser Pro Pro Asp Cys Leu Ser Leu Pro Arg Leu Thr Pro Leu Arg
115 120 125
Leu Pro Pro Pro Pro Phe Pro Phe Leu Gly Pro Ile Pro Ser Leu Pro
130 135 140
Pro Pro Pro Ser Pro Pro Pro Gln Pro Pro Ala Thr Ala Pro Pro Pro
145 150 155 160

```
Ser Leu Arg Phe  *
            164


     <210> 1685
     <211> 153
     <212> PRT
     <213> Homo sapiens


     <400> 1685
Met Gly Thr Ala Ala Leu Gly Pro Val Trp Ala Ala Leu Leu Leu Phe
 1               5                  10                  15
Leu Leu Met Cys Glu Ile Pro Met Val Glu Leu Thr Phe Asp Arg Ala
             20                  25                  30
Val Ala Ser Gly Cys Gln Arg Cys Cys Asp Ser Glu Asp Pro Leu Asp
        35                  40                  45
Pro Ala His Val Ser Ser Ala Ser Ser Ser Gly Arg Pro His Ala Leu
    50                  55                  60
Pro Glu Ile Arg Pro Tyr Ile Asn Ile Thr Ile Leu Lys Ala Gln Arg
 65                  70                  75                  80
Ala Gln His His Ala Glu Pro Glu Cys Asp Ala Gly Pro Gly Leu Arg
                 85                  90                  95
Gly Pro Arg Leu Gly Ala Ala Leu Gln Ala Pro Ala Arg Glu Arg His
            100                 105                 110
Leu Gln Gln Arg Leu Arg His Leu His His Leu Gln Arg Pro Pro His
        115                 120                 125
Gln Gly Arg Gly Arg Leu Arg Ala Ser Gly Pro Pro Ser Arg Leu Glu
    130                 135                 140
Ser Ser Ala Asp Pro Ala Pro Ala  *
145                 150     152


     <210> 1686
     <211> 141
     <212> PRT
     <213> Homo sapiens


     <400> 1686
Met Arg Arg Thr Ala Phe Ile Leu Gly Ser Gly Leu Leu Ser Phe Val
 1               5                  10                  15
Ala Phe Trp Asn Ser Val Thr Trp His Leu Gln Arg Phe Trp Gly Ala
             20                  25                  30
Ser Gly Tyr Phe Trp Gln Ala Gln Trp Glu Arg Leu Leu Thr Thr Phe
        35                  40                  45
Glu Gly Lys Glu Trp Ile Leu Phe Phe Ile Gly Ala Ile Gln Val Pro
    50                  55                  60
Cys Leu Phe Phe Trp Ser Phe Asn Gly Leu Leu Leu Val Val Asp Thr
 65                  70                  75                  80
Thr Gly Lys Pro Asn Phe Ile Ser Arg Tyr Arg Ile Gln Val Gly Lys
                 85                  90                  95
Asn Glu Pro Val Asp Pro Val Lys Leu Arg Gln Ser Ile Arg Thr Val
            100                 105                 110
Leu Phe Asn Gln Cys Met Ile Ser Phe Pro Met Gly Gly Leu Pro Leu
        115                 120                 125
Ser Leu Pro Gln Met Val Glu Arg Pro Leu Thr Pro  *
```

130 135 140

<210> 1687
<211> 61
<212> PRT
<213> Homo sapiens

<400> 1687
Met Leu Thr Glu Leu Leu Leu Leu Cys Val Leu Val Leu Cys Val Phe
1 5 10 15
Met Ser Arg Gly Ser Cys Leu Phe Ala Thr Ile Arg Glu Phe Trp Pro
20 25 30
Pro Trp Val Gly Cys Gly Arg Gly Glu Asn Pro Ser Val Gly Thr Val
35 40 45
Asp Pro Ser Cys Arg Leu Cys Ala Pro Gly His Val *
50 55 60

<210> 1688
<211> 68
<212> PRT
<213> Homo sapiens

<400> 1688
Met Val Ala Ala Thr Pro Pro Gly Ile Ala Arg Trp Ala Leu Val Ile
1 5 10 15
Ser Phe Pro Pro Val Thr Pro Thr Ala Pro His Met Cys Ala Ala Gln
20 25 30
Pro Trp Gly Arg His Gly Ser Ala Glu Gly Thr Thr Gln Leu Pro Ala
35 40 45
Pro Arg Ser Ser Pro Ser Cys Gln Ser Trp Asp Lys Leu Leu Leu Leu
50 55 60
Leu Leu Glu *
65 67

<210> 1689
<211> 74
<212> PRT
<213> Homo sapiens

<400> 1689
Met Ala Ala Thr Met Val Ser Ile Ala Ser Phe Arg Leu Phe Leu Met
1 5 10 15
Ser Cys Thr Leu Val Ala Phe Ser Pro Ser Leu Leu Leu Leu Ala Ala
20 25 30
Cys Gly Ser Ser Ser Pro Pro Ser Pro Leu Asn Pro Leu Thr Cys Arg
35 40 45
Ile Leu Ile Cys Phe Thr Met Val Leu Leu Pro Asp Ser Pro Ala Pro
50 55 60
Ser Ser Ser Arg Arg Cys Val Ala Arg *
65 70 73

<210> 1690
<211> 114
<212> PRT
<213> Homo sapiens

<400> 1690
Met His Met Cys Ala Phe Leu His Val Trp Thr Cys Ala Cys Met His
1               5                   10                  15
Leu Cys Val Cys Val Cys Ala Glu Thr Gly Lys Gly Val Lys Val Leu
            20                  25                  30
Val Arg Glu Pro Gly Ser Phe Leu Phe Pro Asn Leu Ser Cys Ser Lys
        35                  40                  45
Glu Gly Trp Gly Trp Gly Gln Pro Leu Leu Lys Val Ile Gly Glu Glu
    50                  55                  60
Arg Phe Ser Asp Ser Glu Val Thr Ala Ser Val Ala Gln Ala Val Ser
65                  70                  75                  80
Leu Val Thr Val Ile Leu Gln Phe Ser Asp Pro His Val Ser Phe Arg
                85                  90                  95
Gly Lys Arg Lys Lys Gly Thr Leu Trp Trp Val Leu Gly Gly Lys Arg
            100                 105                 110
Lys  *
113

<210> 1691
<211> 69
<212> PRT
<213> Homo sapiens

<400> 1691
Met Ala Phe Leu Leu Ser Thr Leu Leu Asn His Tyr Leu Ala Cys Lys
1               5                   10                  15
His Ser Ser Glu Leu Trp Leu Gln Ser Ser Leu Asn Asn Leu Gly Lys
            20                  25                  30
Lys Lys Asp Lys Ala Tyr Ile Phe Thr Val Leu Ala Leu Lys His Ile
        35                  40                  45
Pro Gln Met Pro Leu Arg Ile Tyr Phe Val Leu Gly Gln Ser Trp Trp
    50                  55                  60
Leu Met Pro Val Ser
65              69

<210> 1692
<211> 103
<212> PRT
<213> Homo sapiens

<400> 1692
Met Leu Gly Pro Thr Val Phe Asn Ile Lys Phe Val Phe Leu Ile Thr
1               5                   10                  15
Ala Leu Gly Ala Leu Pro Ser Ser Leu Pro His Ala His Ser Ala Ala

```
            20                  25                  30
Trp Thr Leu Leu Pro Gly Pro Pro Ala Gln Gln His Ser Thr Arg Leu
        35                  40                  45
Trp Thr Phe Ser Asn Met Ala Gly Val Glu Leu Cys Pro Gly Pro Gln
    50                  55                  60
Pro Ala Gly Pro Ala Ala Pro Val Gly Arg Thr Pro Pro Val Leu Ser
65                  70                  75                  80
Ala Phe Thr Thr Thr Ser Ser Phe Gly Ser Gly Cys Gly Val Thr Ser
                85                  90                  95
Ser Arg Glu Leu Pro Arg Arg
            100         103
```

<210> 1693
<211> 48
<212> PRT
<213> Homo sapiens

<400> 1693

```
Met Gly Arg Phe Leu Asp Glu Gln Trp Val Tyr Phe Ile Ile Leu Leu
1               5                   10                  15
Leu Leu Phe Phe Phe Arg Asp Ser Leu Ala Leu Ser Pro Arg Leu Glu
            20                  25                  30
Cys Ser Gly Ala Ile Ser Val His Ser Lys Leu Arg Leu Pro Gly Ser
        35                  40                  45              48
```

<210> 1694
<211> 92
<212> PRT
<213> Homo sapiens

<400> 1694

```
Met Ile Phe Ala Cys Glu Cys Val Leu Arg Leu Leu Leu Ile Leu Asn
1               5                   10                  15
Val Ser Phe Leu Gly Ala Val Ser Glu Glu Thr Thr Asn Ala Leu Glu
            20                  25                  30
Thr Trp Gly Ala Leu Arg Gln Asp Ile Asn Leu Asp Ile Pro Ser Phe
        35                  40                  45
Leu Leu Arg Glu His Ile Asp Glu Leu Ile Cys Asp Lys Thr Leu Asp
    50                  55                  60
Ser Lys Lys Ile Ala His Phe Arg Ala Glu Lys Glu Thr Phe Ser Glu
65                  70                  75                  80
Lys Asp Thr Tyr Cys Tyr Leu Lys Met Glu Leu  *
                85                  90  91
```

<210> 1695
<211> 83
<212> PRT
<213> Homo sapiens

<400> 1695
Met Ala Val Gln Gln Gln Phe Ile Ile Val Val Leu Arg Leu Val Phe
1 5 10 15
Pro Val Ala Gly Thr Thr Arg Ala Pro Leu His Trp Val Gly Ala Ile
20 25 30
Pro Gly Trp Glu Trp Pro Pro Gly Asp Asp Ala Tyr Pro Ser Leu Leu
35 40 45
Ala Pro Ser Gln His Pro Tyr Ser Gly Glu Ala Leu Cys Leu Leu Leu
50 55 60
Leu Pro Ser Ile Val Leu Leu Glu Ser Cys Arg Lys Val Met Glu Arg
65 70 75 80
Gly Leu *
82

<210> 1696
<211> 159
<212> PRT
<213> Homo sapiens

<400> 1696
Met Leu Trp Leu Phe Gln Ser Leu Leu Phe Val Phe Cys Phe Gly Pro
1 5 10 15
Gly Asn Val Val Ser Gln Ser Ser Leu Thr Pro Leu Met Val Asn Gly
20 25 30
Ile Leu Gly Glu Ser Val Thr Leu Pro Leu Glu Phe Pro Ala Gly Glu
35 40 45
Lys Val Asn Phe Ile Thr Trp Leu Phe Asn Glu Thr Ser Leu Ala Phe
50 55 60
Ile Val Pro His Glu Thr Lys Ser Pro Glu Ile His Val Thr Asn Pro
65 70 75 80
Lys Gln Gly Lys Arg Leu Asn Phe Thr Gln Ser Tyr Ser Leu Gln Leu
85 90 95
Ser Asn Leu Lys Met Glu Asp Thr Gly Ser Tyr Arg Ala Gln Ile Ser
100 105 110
Thr Lys Thr Ser Ala Lys Leu Ser Ser Tyr Thr Leu Arg Ile Leu Thr
115 120 125
Leu Tyr Pro Ile Val Gly Asn Gly Ile Trp Gly Asn Lys Asn Phe Leu
130 135 140
Thr Thr Leu Ala Arg Gly Asn Val Lys Leu Asp Gly Leu His Glu
145 150 155 159

<210> 1697
<211> 105
<212> PRT
<213> Homo sapiens

<400> 1697
Met Glu Pro Arg Leu Phe Cys Trp Thr Thr Leu Phe Leu Leu Ala Gly
1 5 10 15
Trp Cys Leu Pro Gly Leu Pro Cys Pro Ser Arg Cys Leu Cys Phe Lys
20 25 30
Ser Thr Val Arg Cys Met His Leu Met Leu Asp His Ile Pro Gln Val

```
        35                  40                  45
Pro Gln Gln Thr Thr Val Leu Asp Leu Arg Phe Asn Arg Ile Arg Glu
    50                  55                  60
Ile Pro Gly Ser Ala Phe Lys Lys Leu Lys Asn Leu Asn Thr Leu Tyr
65                  70                  75                  80
Leu Tyr Lys Asn Glu Ile His Ala Leu Asp Lys Gln Thr Phe Lys Gly
                85                  90                  95
Leu Ile Ser Leu Glu His Leu Tyr Ile
            100                 105
```

<210> 1698
<211> 195
<212> PRT
<213> Homo sapiens

<400> 1698

```
Met Pro Ser Trp Ile Gly Ala Val Ile Leu Pro Leu Leu Gly Leu Leu
 1               5                  10                  15
Leu Ser Leu Pro Ala Gly Ala Asp Val Lys Ala Arg Ser Cys Gly Glu
            20                  25                  30
Val Arg Gln Ala Tyr Gly Ala Lys Gly Phe Ser Leu Ala Asp Ile Pro
        35                  40                  45
Tyr Gln Glu Ile Ala Gly Glu His Leu Arg Ile Cys Pro Gln Glu Tyr
    50                  55                  60
Thr Cys Cys Thr Thr Glu Met Glu Asp Lys Leu Ser Gln Gln Ser Lys
65                  70                  75                  80
Leu Glu Phe Glu Asn Leu Val Glu Glu Thr Ser His Phe Val Arg Thr
                85                  90                  95
Thr Phe Val Ser Arg His Lys Lys Phe Asp Glu Phe Phe Arg Glu Leu
            100                 105                 110
Leu Glu Asn Ala Glu Lys Ser Leu Asn Asp Met Phe Val Arg Thr Tyr
        115                 120                 125
Gly Met Leu Tyr Met Gln Asn Ser Glu Val Phe Gln Asp Leu Phe Thr
    130                 135                 140
Glu Leu Lys Arg Tyr Tyr Thr Gly Gly Asn Val Asn Leu Glu Glu Met
145                 150                 155                 160
Leu Asn Asp Phe Trp Ala Arg Leu Leu Glu Arg Met Phe Gln Leu Ile
                165                 170                 175
Asn Pro Gln Tyr Pro Phe Ser Glu Gly Phe Leu Gly Met Cys Glu Gln
            180                 185                 190
Ile Pro  *
    194
```

<210> 1699
<211> 97
<212> PRT
<213> Homo sapiens

<400> 1699

```
Met Asp Ser Pro Trp Ala Gly Leu Leu Trp Leu Leu Pro Thr Leu Trp
 1               5                  10                  15
Ser Ser Phe Pro Ala Pro Ala Cys Trp Pro Ser Ser Ser Ser Ser Ser
            20                  25                  30
```

Pro Val Cys Ala Ala Asn Gly Ala Met Ser Ala Ser Arg Asn Leu Arg
35 40 45
Thr Leu Lys Gly Arg Thr Ala Pro Gly Ser Thr Leu Pro Leu Arg Arg
50 55 60
Arg Pro Pro Pro His Ser Arg Cys Leu Met Ser Thr Phe Ser Arg Trp
65 70 75 80
Leu Arg Ser Pro Cys Gln Cys Leu Pro Arg Ser Leu His Thr Gln Thr
85 90 95 96
*

<210> 1700
<211> 129
<212> PRT
<213> Homo sapiens

<400> 1700
Met Gly Trp Ala Pro Leu Leu Leu Thr Leu Leu Ala His Cys Thr Gly
1 5 10 15
Ser Trp Ala Gln Ser Val Leu Thr Gln Pro Pro Ser Glu Ser Glu Ala
20 25 30
Pro Gly Gln Trp Val Asn Ile Ser Cys Thr Gly Ser Gly Ser Asn Leu
35 40 45
Gly Ala Gly Phe Asp Val Gln Trp Tyr Gln Leu Ile Pro Gly Thr Ala
50 55 60
Pro Lys Leu Leu Ile Phe Asn Asn Asn Arg Gln Pro Ser Gly Val Pro
65 70 75 80
Asp Arg Phe Ser Ala Ser Lys Ser Gly Thr Ser Ala Ser Leu Thr Ile
85 90 95
Asn Asp Leu Gln Pro Glu Asp Glu Ser Glu Tyr Tyr Cys Leu Ala Met
100 105 110
Thr Ala Ala Ser Leu Val Ser Ser Glu Leu Gly Pro Lys Ser Pro Ala
115 120 125 128
*

<210> 1701
<211> 219
<212> PRT
<213> Homo sapiens

<400> 1701
Met Arg Thr His Thr Arg Gly Ala Pro Ser Val Phe Phe Ile Tyr Leu
1 5 10 15
Leu Cys Phe Val Ser Ala Tyr Ile Thr Asp Glu Asn Pro Glu Val Met
20 25 30
Ile Pro Phe Thr Asn Ala Asn Tyr Asp Ser His Pro Met Leu Tyr Phe
35 40 45
Ser Arg Ala Glu Val Ala Glu Leu Gln Leu Arg Ala Ala Ser Ser His
50 55 60
Glu His Ile Ala Ala Arg Leu Thr Glu Ala Val His Thr Met Leu Ser
65 70 75 80
Ser Pro Leu Glu Tyr Leu Pro Pro Trp Asp Pro Lys Asp Tyr Ser Ala

```
                85                  90                  95
Arg Trp Asn Glu Ile Phe Gly Asn Asn Leu Gly Ala Leu Ala Met Phe
            100                 105                 110
Cys Val Leu Tyr Pro Glu Asn Ile Glu Ala Arg Asp Met Ala Lys Asp
        115                 120                 125
Tyr Met Glu Arg Met Ala Ala Gln Pro Ser Trp Leu Val Lys Asp Ala
    130                 135                 140
Pro Trp Asp Glu Val Pro Leu Ala His Ser Leu Val Gly Phe Ala Thr
145                 150                 155                 160
Ala Tyr Asp Phe Leu Tyr Asn His Leu Ser Lys Thr Gln Gln Glu Lys
                165                 170                 175
Phe Leu Glu Val Ile Ala Asn Ala Ser Gly Tyr Met Phe Val Thr Leu
            180                 185                 190
Ile Leu Gly Ala Asp Gly Asp Ser Asn Thr Cys Thr Ile Ile Ser Pro
        195                 200                 205
Pro Thr Val Trp Leu Cys Ser Arg Glu Ala  *
    210                 215         218
```

<210> 1702
<211> 86
<212> PRT
<213> Homo sapiens

<400> 1702

```
Met Glu Gln Leu Leu Gly Ile Lys Leu Gly Cys Leu Phe Ala Leu Leu
1               5                   10                  15
Ala Leu Thr Leu Gly Cys Gly Leu Thr Pro Ile Cys Phe Lys Trp Phe
            20                  25                  30
Gln Ile Asp Ala Ala Arg Gly His His Arg Leu Val Leu Arg Leu Leu
        35                  40                  45
Gly Cys Ile Ser Ala Gly Val Phe Leu Gly Ala Gly Phe Met His Met
    50                  55                  60
Thr Ala Glu Ala Leu Glu Glu Ile Glu Ser Gln Ile Gln Lys Phe Met
65                  70                  75                  80
Val Gln Ile Ser Lys  *
                85
```

<210> 1703
<211> 229
<212> PRT
<213> Homo sapiens

<400> 1703

```
Met Leu Ser Met Leu Arg Thr Met Thr Arg Leu Cys Phe Leu Leu Phe
1               5                   10                  15
Phe Ser Val Ala Thr Ser Gly Cys Ser Ala Ala Ala Ala Ser Ser Leu
            20                  25                  30
Glu Met Leu Ser Arg Glu Phe Glu Thr Cys Ala Phe Ser Phe Ser Ser
        35                  40                  45
Leu Pro Arg Ser Cys Lys Glu Ile Lys Glu Arg Cys His Ser Ala Gly
    50                  55                  60
Asp Gly Leu Tyr Phe Leu Arg Thr Lys Asn Gly Val Val Tyr Gln Thr
65                  70                  75                  80
```

Phe Cys Asp Met Thr Ser Gly Gly Gly Gly Trp Thr Leu Val Ala Ser
85 90 95
Val His Glu Asn Asp Met His Gly Lys Cys Thr Val Gly Asp Arg Trp
100 105 110
Ser Ser Gln Gln Gly Asn Lys Ala Asp Tyr Pro Glu Gly Asp Gly Asn
115 120 125
Trp Ala Asn Tyr Asn Thr Phe Gly Ser Ala Glu Ala Ala Thr Ser Asp
130 135 140
Asp Tyr Lys Asn Pro Gly Tyr Tyr Asp Ile Gln Ala Lys Asp Leu Gly
145 150 155 160
Ile Trp His Val Pro Asn Lys Ser Pro Met Gln His Trp Arg Asn Ser
165 170 175
Ala Leu Leu Arg Tyr Arg Thr Asn Thr Gly Phe Leu Gln Arg Leu Gly
180 185 190
His Asn Leu Phe Gly Ile Tyr Gln Lys Tyr Pro Val Lys Tyr Arg Ser
195 200 205
Gly Lys Cys Trp Asn Asp Asn Gly Pro Ala Ile Pro Trp Val Tyr Asp
210 215 220
Phe Gly Glu Ala *
225 228

<210> 1704
<211> 202
<212> PRT
<213> Homo sapiens

<400> 1704
Met Val Phe Pro Val Met Tyr Asn Leu Ile Ile Leu Val Cys Arg Ala
1 5 10 15
Cys Phe Pro Asp Leu Gln His Gly Tyr Leu Val Ala Trp Leu Val Leu
20 25 30
Asp Tyr Thr Ser Asp Leu Leu Tyr Leu Leu Asp Met Val Val Arg Phe
35 40 45
His Thr Gly Phe Leu Glu Gln Gly Ile Leu Val Val Asp Lys Gly Arg
50 55 60
Ile Ser Ser Arg Tyr Val Arg Thr Trp Ser Phe Phe Leu Asp Leu Ala
65 70 75 80
Ser Leu Met Pro Thr Asp Val Val Tyr Val Arg Leu Gly Pro His Thr
85 90 95
Pro Thr Leu Arg Leu Asn Arg Phe Leu Arg Ala Pro Arg Leu Phe Glu
100 105 110
Ala Phe Asp Arg Thr Glu Thr Arg Thr Ala Tyr Pro Asn Ala Phe Cys
115 120 125
Ile Gly Lys Leu Met Leu Tyr Ile Phe Gly Arg Ile His Trp Asn Asn
130 135 140
Cys Leu Tyr Phe Ser Leu Ser Arg Tyr Leu Gly Phe Gly Arg Glu Pro
145 150 155 160
Met Gly Val Pro Arg Thr Pro Ala Pro Thr Trp Val Leu Thr Ala Arg
165 170 175
Gly Gly Pro Val Thr Ser Tyr Lys Leu Phe Asn Phe Phe His Pro Leu
180 185 190
Asp Thr Trp Ile Ile Gln Gly Gly Glu *
195 200 201

<210> 1705
<211> 58
<212> PRT
<213> Homo sapiens

<400> 1705
Met Gly Leu Leu Gly Val Leu Trp Asn Thr Thr Leu His Met Cys Arg
1 5 10 15
Met Arg Leu Gln Asp Thr Gly Gln Lys Ile Arg Thr Gly Ser Cys Glu
20 25 30
Leu His Gly Ser Gln Ser Ser His Ser Thr Gly Asn Leu Arg Val Leu
35 40 45
Pro Ser His Asn Gly Glu Thr Leu His *
50 55 57

<210> 1706
<211> 55
<212> PRT
<213> Homo sapiens

<400> 1706
Met Gly Asp Tyr Arg Asn Val Arg Leu Leu Gly Ser Phe Ser Phe Ile
1 5 10 15
Ser Val Thr Ile Ser Arg Val Ile Phe Leu Leu Ser Leu Leu Gln Pro
20 25 30
Ser Gly Val Gly Ile Leu Phe Ala Asp Ser Gly Gly Thr Gly Tyr Thr
35 40 45
His His Cys Leu Trp Val *
50 54

<210> 1707
<211> 139
<212> PRT
<213> Homo sapiens

<400> 1707
Met Leu Glu Cys Ala Phe Ile Val Leu Trp Leu Gln Leu Gly Trp Leu
1 5 10 15
Ser Gly Glu Asp Gln Val Thr Gln Ser Pro Glu Ala Leu Arg Leu Gln
20 25 30
Glu Gly Glu Ser Ser Ser Leu Asn Cys Ser Tyr Thr Val Ser Gly Leu
35 40 45
Arg Gly Leu Phe Trp Tyr Arg Gln Asp Pro Gly Lys Gly Pro Glu Phe
50 55 60
Leu Phe Thr Leu Tyr Ser Ala Gly Glu Glu Lys Glu Lys Glu Arg Leu
65 70 75 80
Lys Ala Thr Leu Thr Lys Lys Glu Ser Phe Leu His Ile Thr Ala Pro
85 90 95
Lys Pro Glu Asp Ser Ala Thr Tyr Leu Cys Ala Val Gln Ala Gln Phe
100 105 110
His Ser Gly Gly Gly Ala Asp Gly Leu Thr Phe Gly Lys Gly Thr Arg
115 120 125

```
Leu Lys Val Leu Ala Leu Tyr Pro Glu Pro  *
    130                 135         138



     <210> 1708
     <211> 59
     <212> PRT
     <213> Homo sapiens


     <400> 1708
Met Gly Pro Arg Phe Val Ser Thr Leu Pro Phe Ser Pro Ser Ala Ala
 1               5                  10                  15
Trp Cys Ala Cys Glu Ala Gly Gly Gly Leu Arg Arg Glu Val Ala His
            20                  25                  30
Ala Gln Arg Ala Ala Ser Thr Ala Pro Thr Ala His Met Gln Asn Ser
        35                  40                  45
Thr Leu Ile Gly Leu Asn Leu Ser Arg Gly  *
     50                  55         58



     <210> 1709
     <211> 81
     <212> PRT
     <213> Homo sapiens


     <400> 1709
Met Arg Leu Pro Trp Glu Leu Leu Val Leu Gln Ser Phe Ile Leu Cys
 1               5                  10                  15
Leu Ala Asp Asp Ser Thr Leu His Gly Pro Ile Phe Ile Gln Glu Pro
            20                  25                  30
Ser Pro Val Met Phe Pro Leu Asp Ser Glu Glu Lys Lys Ala Lys Leu
        35                  40                  45
Asn Cys Glu Asp Lys Gly Asp Pro Lys Pro His Ile Arg Trp Lys Leu
     50                  55                  60
Asn Gly Ala Asp Ala Asp Thr Gly Met Glu Phe Leu Leu Gln Arg Cys
65                  70                  75                  80
 *



     <210> 1710
     <211> 399
     <212> PRT
     <213> Homo sapiens


     <400> 1710
Met Leu Arg Leu Tyr Val Leu Val Met Gly Val Ser Ala Phe Thr Leu
 1               5                  10                  15
Gln Pro Ala Ala His Thr Gly Ala Ala Arg Ser Cys Arg Phe Arg Gly
            20                  25                  30
Arg His Tyr Lys Arg Glu Phe Arg Leu Glu Gly Glu Pro Val Ala Leu
        35                  40                  45
Arg Cys Pro Gln Val Pro Tyr Trp Leu Trp Ala Ser Val Ser Pro Arg
```

```
 50                  55                  60
Ile Asn Leu Thr Trp His Lys Asn Asp Ser Ala Arg Thr Val Pro Gly
 65                  70                  75                  80
Glu Glu Glu Thr Arg Met Trp Ala Gln Asp Gly Ala Leu Trp Leu Leu
                 85                  90                  95
Pro Ala Leu Gln Glu Asp Ser Gly Thr Tyr Val Cys Thr Thr Arg Asn
            100                 105                 110
Ala Ser Tyr Cys Asp Lys Met Ser Ile Glu Leu Arg Val Phe Glu Asn
        115                 120                 125
Thr Asp Ala Phe Leu Pro Phe Ile Ser Tyr Pro Gln Ile Leu Thr Leu
    130                 135                 140
Ser Thr Ser Gly Val Leu Val Cys Pro Asp Leu Ser Glu Phe Thr Arg
145                 150                 155                 160
Asp Lys Thr Asp Val Lys Ile Gln Trp Tyr Lys Asp Ser Leu Leu Leu
                165                 170                 175
Asp Lys Asp Asn Glu Lys Phe Leu Ser Val Arg Gly Thr Thr His Leu
            180                 185                 190
Leu Val His Asp Val Ala Leu Glu Asp Ala Gly Tyr Tyr Arg Cys Val
        195                 200                 205
Leu Thr Phe Ala His Glu Gly Gln Gln Tyr Asn Ile Thr Arg Ser Ile
    210                 215                 220
Glu Leu Arg Ile Lys Lys Lys Lys Glu Glu Thr Ile Pro Val Ile Ile
225                 230                 235                 240
Ser Pro Leu Lys Thr Ile Ser Ala Ser Leu Gly Ser Arg Leu Thr Ile
                245                 250                 255
Pro Cys Lys Val Phe Leu Gly Thr Gly Thr Pro Leu Thr Thr Met Leu
            260                 265                 270
Trp Trp Thr Ala Asn Asp Thr His Ile Glu Ser Ala Tyr Pro Gly Gly
        275                 280                 285
Arg Val Thr Glu Gly Pro Arg Gln Glu Tyr Ser Glu Asn Asn Glu Asn
    290                 295                 300
Tyr Ile Glu Val Pro Leu Ile Phe Asp Pro Val Thr Arg Glu Asp Leu
305                 310                 315                 320
His Met Asp Phe Lys Cys Val Val His Asn Thr Leu Ser Phe Gln Thr
                325                 330                 335
Leu Arg Thr Thr Val Lys Glu Ala Ser Ser Thr Phe Ser Trp Gly Ile
            340                 345                 350
Val Leu Ala Pro Leu Ser Leu Ala Phe Leu Val Leu Gly Gly Ile Trp
        355                 360                 365
Met His Arg Arg Cys Lys His Arg Thr Gly Lys Ala Asp Gly Leu Thr
    370                 375                 380
Val Leu Trp Pro His His Gln Asp Phe Gln Ser Tyr Pro Lys  *
385                 390                 395         398
```

```
<210> 1711
<211> 254
<212> PRT
<213> Homo sapiens

<400> 1711
Met Ala Met Gly Val Pro Arg Val Ile Leu Leu Cys Leu Phe Gly Ala
 1               5                  10                  15
Ala Leu Cys Leu Thr Gly Ser Gln Ala Leu Gln Cys Tyr Ser Phe Glu
            20                  25                  30
His Thr Tyr Phe Gly Pro Phe Asp Leu Arg Ala Met Lys Leu Pro Ser
        35                  40                  45
```

Ile Ser Cys Pro His Glu Cys Phe Glu Ala Ile Leu Ser Leu Asp Thr
50 55 60
Gly Tyr Arg Ala Pro Val Thr Leu Val Arg Lys Gly Cys Trp Thr Gly
65 70 75 80
Pro Pro Ala Gly Gln Thr Gln Ser Asn Ala Asp Ala Leu Pro Pro Asp
85 90 95
Tyr Ser Val Val Arg Gly Cys Thr Thr Asp Lys Cys Asn Ala His Leu
100 105 110
Met Thr His Asp Ala Leu Pro Asn Leu Ser Gln Ala Pro Asp Pro Pro
115 120 125
Thr Leu Ser Gly Leu Glu Cys Tyr Ala Cys Ile Gly Val His Gln Asp
130 135 140
Asp Cys Ala Ile Gly Arg Ser Arg Arg Val Gln Cys His Gln Asp Gln
145 150 155 160
Thr Ala Cys Phe Gln Gly Asn Gly Arg Met Thr Val Gly Asn Phe Ser
165 170 175
Val Pro Val Tyr Ile Arg Thr Cys His Arg Ala Leu Leu His His Leu
180 185 190
Met Gly Thr Thr Ser Pro Trp Thr Ala Ile Gly Pro Pro Arg Gly Ser
195 200 205
Cys Cys Glu Gly Tyr Leu Cys Asn Arg Lys Ser Met Thr Gln Pro Phe
210 215 220
Thr Ser Ala Ser Ala Thr Thr Pro Pro Arg Ala Leu Gln Val Leu Ala
225 230 235 240
Leu Leu Leu Pro Val Leu Leu Leu Val Gly Leu Ser Ala *
245 250 253

<210> 1712
<211> 124
<212> PRT
<213> Homo sapiens

<400> 1712
Met Thr Trp Leu Leu Val Ala Tyr Ala Asp Phe Val Val Thr Phe Val
1 5 10 15
Met Leu Leu Pro Ser Lys Asp Phe Trp Tyr Ser Val Val Asn Gly Val
20 25 30
Ile Phe Asn Cys Leu Ala Val Leu Ala Leu Ser Ser His Leu Arg Thr
35 40 45
Met Leu Thr Asp Pro Glu Lys Ser Ser Asp Cys Arg Pro Ser Ala Cys
50 55 60
Thr Val Lys Thr Gly Leu Asp Pro Thr Leu Val Gly Ile Cys Gly Glu
65 70 75 80
Gly Thr Glu Ser Val Gln Ser Leu Leu Leu Gly Ala Val Pro Lys Gly
85 90 95
Asn Ala Thr Lys Glu Tyr Met Asp Glu Leu Ala Ala Glu Ala Arg Gly
100 105 110
Ser His Leu Gln Val Pro Gln Val Leu Leu Tyr *
115 120 123

<210> 1713
<211> 214
<212> PRT
<213> Homo sapiens

<400> 1713
Met Leu His Leu Val Phe Ile Leu Pro Ser Leu Met Leu Leu Ile Pro
1 5 10 15
His Ile Leu Leu Glu Asn Phe Ala Ala Ala Ile Pro Gly His Arg Cys
20 25 30
Trp Val His Met Leu Asp Asn Asn Thr Gly Ser Gly Asn Glu Thr Gly
35 40 45
Ile Leu Ser Glu Asp Ala Leu Leu Arg Ile Ser Ile Pro Leu Asp Ser
50 55 60
Asn Leu Arg Pro Glu Lys Cys Arg Arg Phe Val His Pro Gln Trp Gln
65 70 75 80
Leu Leu His Leu Asn Gly Thr Ile His Ser Thr Ser Glu Ala Asp Thr
85 90 95
Glu Pro Cys Val Asp Gly Trp Val Tyr Asp Gln Ser Tyr Phe Pro Ser
100 105 110
Thr Ile Val Thr Lys Trp Asp Leu Val Cys Asp Tyr Gln Ser Leu Lys
115 120 125
Ser Val Val Gln Phe Leu Leu Leu Thr Gly Met Leu Val Gly Gly Ile
130 135 140
Ile Gly Gly His Val Ser Asp Arg Trp Leu Val Glu Ser Ala Arg Trp
145 150 155 160
Leu Ile Ile Thr Asn Lys Leu Asp Glu Gly Leu Lys Ala Leu Arg Lys
165 170 175
Val Ala Arg Thr Asn Gly Ile Lys Asn Ala Glu Arg Asn Pro Glu His
180 185 190
Arg Gly Cys Lys Ile His His Ala Gly Gly Ala Gly Cys Ser Thr Asp
195 200 205
Gln Asn Tyr Cys Val *
210 213

<210> 1714
<211> 178
<212> PRT
<213> Homo sapiens

<400> 1714
Met Ala Ala Ser Trp Ser Leu Leu Val Thr Leu Arg Pro Leu Ala Gln
1 5 10 15
Ser Pro Leu Arg Gly Arg Cys Val Gly Cys Gly Ala Trp Ala Ala Ala
20 25 30
Leu Ala Pro Leu Ala Thr Ala Pro Gly Lys Pro Phe Trp Lys Ala Tyr
35 40 45
Thr Val Gln Thr Ser Glu Ser Met Thr Pro Thr Ala Thr Ser Glu Thr
50 55 60
Tyr Leu Lys Ala Leu Ala Val Cys His Gly Pro Leu Asp His Tyr Asp
65 70 75 80
Phe Leu Ile Lys Ala His Glu Leu Lys Asp Asp Glu His Gln Arg Arg
85 90 95
Val Ile Gln Cys Leu Gln Lys Leu His Glu Asp Leu Lys Gly Tyr Asn
100 105 110
Ile Glu Ala Glu Gly Leu Phe Phe Lys Ala Phe Phe Lys Glu Gln Thr
115 120 125
Ser Lys Gly Pro Val Cys Leu Trp Arg Cys Trp Tyr Arg Lys Asn Asn
130 135 140

Gly Asp Gly His Val Leu Cys Leu Cys Gly Asn Glu Glu Glu Lys Thr
145 150 155 160
Gly Ser Phe Ser Trp Phe His Ala Arg Cys Ala Gln Lys Asn Thr Ser
165 170 175
Pro *
177

<210> 1715
<211> 76
<212> PRT
<213> Homo sapiens

<400> 1715
Met Arg Val Thr Ala Pro Arg Thr Val Leu Leu Leu Leu Trp Gly Ala
1 5 10 15
Val Ala Leu Thr Glu Thr Trp Ala Gly Ser His Ser Met Lys Tyr Phe
20 25 30
Tyr Thr Ala Met Ser Arg Ala Gly Arg Gly Glu Pro Arg Phe Ile Ala
35 40 45
Glu Gly Tyr Val Asp Asp Thr Gln Phe Val Arg Phe Asp Ser Asp Ala
50 55 60
Ala Ser Pro Lys Thr Asp Pro Gly Arg His Gly *
65 70 75

<210> 1716
<211> 83
<212> PRT
<213> Homo sapiens

<400> 1716
Met Arg Phe Thr Phe Pro Leu Met Ala Ile Val Leu Glu Ile Ala Met
1 5 10 15
Ile Ala Ser Phe Gly Leu Phe Val Glu Tyr Glu Thr Asp His Thr Val
20 25 30
Leu Glu His Phe Asn Ile Thr Lys Pro Ser Asp Met Gly Ile Phe Phe
35 40 45
Glu Leu Tyr Pro Leu Phe Gln Asp Val His Gly Met Ile Phe Val Gly
50 55 60
Phe Asp Phe Pro Pro Asp Leu Pro Glu Glu Leu Trp Val Ser Gln Arg
65 70 75 80
Gly Tyr *
82

<210> 1717
<211> 57
<212> PRT
<213> Homo sapiens

<400> 1717
Met Ala Leu Phe Phe Leu Ala Leu Asn Phe Trp Lys Val Gly Met Ala

```
 1               5                   10                  15
Cys Tyr Val Arg Thr Ser Ser Trp Asn Ser Leu Leu Phe Phe Ser Gln
            20                  25                  30
Pro Tyr Phe Leu Gly Ser Cys Phe Glu Gln Tyr Leu Ser Asn Val Cys
        35                  40                  45
Leu Pro Asp Val Val Pro Asp Ala  *
    50                  55  56
```

<210> 1718
<211> 76
<212> PRT
<213> Homo sapiens

<400> 1718

```
Met Tyr Leu Gly Leu Phe Leu Asp Phe Tyr Ser Val Ser Phe Cys Gly
 1               5                  10                  15
Cys Leu His Met Leu Gln Pro Gln Cys Phe Asn Tyr Phe Asn Ser Lys
            20                  25                  30
Asp Gln Ser Arg Phe His Cys Leu Lys His Cys Ser Asp His Leu Ile
        35                  40                  45
Phe Leu Leu Ser Glu Leu Arg Ser Asn Met Phe Ser Ser Phe Leu Ile
    50                  55                  60
Leu Ser Ile Phe Tyr Asp Tyr Cys Ile Asn Leu  *
65                  70                  75
```

<210> 1719
<211> 71
<212> PRT
<213> Homo sapiens

<400> 1719

```
Met Lys Ile Phe Phe His Ile Phe Phe His Lys Cys Leu Phe Thr Tyr
 1               5                  10                  15
Arg Leu Phe Ile Thr Leu Ala Leu Ile Leu Trp Tyr Ser Asp Ile Glu
            20                  25                  30
Glu Ser Thr Phe Pro Pro Leu Met Arg Tyr Cys Pro Asn Thr Val Leu
        35                  40                  45
His Lys Ser Phe Phe Gln Met Ser Ala Phe Ile Thr Tyr Gln Phe Ser
    50                  55                  60
Leu Tyr Leu Ser Leu Phe  *
65                  70
```

<210> 1720
<211> 101
<212> PRT
<213> Homo sapiens

<400> 1720

```
Met Leu Ala Gly Gln Leu Leu Pro Met Leu Thr Leu Leu Pro Pro Ser
 1               5                  10                  15
```

```
Phe Pro Leu Pro His Pro Thr Leu Gly Pro Arg Arg His Ala Ser Leu
            20                  25                  30
Thr Gln Leu Gly Pro Ala Phe Trp Met Ala Trp Gly Arg Pro Trp Ala
        35                  40                  45
His Leu Gly Pro Gly Gln Pro Leu Gly Gln Leu Trp Lys Ser Ser Val
    50                  55                  60
Glu Glu His Leu Leu Ala Ala Trp Leu Gln Pro Leu Ala Leu Leu Glu
65                  70                  75                  80
Trp Ser Leu Gly Ala Ser Ala Leu Ser Ala Leu Gly Thr Ser His Pro
                85                  90                  95
Leu Gly Leu Gln  *
            100
```

```
     <210> 1721
     <211> 48
     <212> PRT
     <213> Homo sapiens
```

```
     <400> 1721
Met Leu Val Leu Leu Val Trp Val His His Thr Leu Leu Leu Gly Gln
 1               5                  10                  15
Lys Ser Thr Tyr Glu Glu Lys Arg Asn Gly Lys Trp Gly Arg Gln Arg
            20                  25                  30
Arg Ala Pro Tyr Leu Gly Val Tyr Ile Glu Ala Thr Gly Gln Val  *
        35                  40                  45      47
```

```
     <210> 1722
     <211> 70
     <212> PRT
     <213> Homo sapiens
```

```
     <400> 1722
Met Asp Val Gly Pro Asn Ser Leu Pro His Leu Gly Leu Lys Leu Leu
 1               5                  10                  15
Leu Leu Leu Leu Leu Val Thr Leu Arg Gly Gln Ala Asn Thr Gly Trp
            20                  25                  30
Tyr Gly Ile Pro Gly Met Pro Gly Leu Pro Gly Ala Pro Gly Lys Asp
        35                  40                  45
Gly Tyr Asp Gly Leu Pro Gly Pro Lys Gly Glu Pro Gly Ile Asp Ala
    50                  55                  60
Ile Ser Leu Ile Leu  *
65              69
```

```
     <210> 1723
     <211> 54
     <212> PRT
     <213> Homo sapiens
```

```
     <400> 1723
Met Asp Leu Ile Phe Val Lys Val Leu Leu Ile Phe Ala Ala Ile Gln
```

```
 1               5                   10                  15
Thr Leu Ser Lys Trp Gln Phe Ala Phe Thr Phe Ser Ile Gln Thr Val
            20                  25                  30
Pro Ser Leu Val Ile Asn Leu Ser Trp Leu Leu Leu Asp Leu Lys Pro
        35                  40                  45
Gly Thr His Ile Gln  *
    50          53
```

<210> 1724
<211> 60
<212> PRT
<213> Homo sapiens

<400> 1724

```
Met Val Ser Gly Trp Ile Thr Lys Thr Gln Phe Leu Leu Leu Gly Arg
 1               5                   10                  15
Gly Lys Ile Cys Met Tyr Lys Cys Ile Lys Gln Leu Gln Val Arg Lys
            20                  25                  30
Thr Asp Val Ile Thr Thr Lys Gln Ile Asn Tyr Glu Glu Ile Asn Cys
        35                  40                  45
Leu Asn His Ile Met Leu Thr Thr Lys Phe Trp  *
    50                  55              59
```

<210> 1725
<211> 63
<212> PRT
<213> Homo sapiens

<400> 1725

```
Met Phe Phe Arg Met Gln Val Cys Glu His His Gly Phe Trp Val Ile
 1               5                   10                  15
Leu Leu Leu Leu Ser Leu Lys Met Glu Ile Pro Leu Ala Ala Tyr Pro
            20                  25                  30
Thr Ala Glu Tyr Ser Ser Ile Gly Ser Gly Phe Thr Pro Leu His Pro
        35                  40                  45
Ser Arg Thr Phe Thr Gln Ala Ser Pro Leu Pro Ser Ile Phe  *
    50                  55                  60      62
```

<210> 1726
<211> 57
<212> PRT
<213> Homo sapiens

<400> 1726

```
Met Cys Leu Phe Cys Ser Phe Val Asn Val Thr Leu Gly Ser Thr Asp
 1               5                   10                  15
Pro Met Cys Cys Pro Ala Gln Trp Leu Ala Gln Arg Met Pro Trp Ala
            20                  25                  30
Phe Val Ser Ile Arg Lys Ala Trp Pro Leu Gly Arg Met Ser Gly Ala
        35                  40                  45
```

Ser Gln Arg Leu Lys Glu Glu Glu *
50 55 56

<210> 1727
<211> 46
<212> PRT
<213> Homo sapiens

<400> 1727
Met Arg Trp Pro Trp Ala Ser Trp Ala Ala Val Leu Leu Lys Leu Pro
1 5 10 15
Arg Arg Val Leu Pro Trp Leu Pro Cys Gly His Gln Gln His Val Arg
20 25 30
Ala Thr Ala Ser Ser Arg Ser Pro Pro Met Pro Val Thr Lys
35 40 45 46

<210> 1728
<211> 46
<212> PRT
<213> Homo sapiens

<400> 1728
Met Lys Met Glu Met Glu Thr Lys Arg Ser Trp Arg Pro Gln Ser His
1 5 10 15
Gly His Phe Thr Phe Gln Phe Leu Leu Ser Trp Thr Phe Glu Leu Ile
20 25 30
Leu Phe His Phe Val Pro Phe Phe Pro Tyr Leu Leu Phe *
35 40 45

<210> 1729
<211> 49
<212> PRT
<213> Homo sapiens

<400> 1729
Met Val Leu Leu Pro Leu Gln Cys Gly Leu Thr Lys Ala Ser Ser Cys
1 5 10 15
Leu His Thr Leu Cys Ser Ser Ser Asp Gln Ile Gly Tyr Leu Pro Val
20 25 30
Lys Asn Thr Asp Gln Leu Gly Leu Gln Met Glu Val Ala Glu Met Cys
35 40 45 48
*

<210> 1730
<211> 50
<212> PRT
<213> Homo sapiens

<400> 1730
Met Phe Thr Phe Gly Arg Leu Phe Gln Ile Ile Thr Val Val Thr Cys
1 5 10 15
Leu Gln Phe Ile Gln Asp Cys Cys Ile His Ser Arg Gln Ile Asn Ser
20 25 30
Leu Leu Glu Thr Ser Ser Leu Ser Arg Cys Leu Glu Val Pro Asp Val
35 40 45
Cys *
49

<210> 1731
<211> 227
<212> PRT
<213> Homo sapiens

<400> 1731
Met Gly Cys Asp Gly Arg Val Ser Gly Leu Leu Arg Arg Asn Leu Gln
1 5 10 15
Pro Thr Leu Thr Tyr Trp Ser Val Phe Phe Ser Phe Gly Leu Cys Ile
20 25 30
Ala Phe Leu Gly Pro Thr Leu Leu Asp Leu Arg Cys Gln Thr His Ser
35 40 45
Ser Leu Pro Gln Ile Ser Trp Val Phe Phe Ser Gln Gln Leu Cys Leu
50 55 60
Leu Leu Gly Ser Ala Leu Gly Gly Val Phe Lys Arg Thr Leu Ala Gln
65 70 75 80
Ser Leu Trp Ala Leu Phe Thr Ser Ser Leu Ala Ile Ser Leu Val Phe
85 90 95
Ala Val Ile Pro Phe Cys Arg Asp Val Lys Val Leu Ala Ser Val Met
100 105 110
Ala Leu Ala Gly Leu Ala Met Gly Cys Ile Asp Thr Val Ala Asn Met
115 120 125
Gln Leu Val Arg Met Tyr Gln Lys Asp Ser Ala Val Phe Leu Gln Val
130 135 140
Leu His Phe Phe Val Gly Phe Gly Ala Leu Leu Ser Pro Leu Ile Ala
145 150 155 160
Asp Pro Phe Leu Ser Glu Ala Asn Cys Leu Pro Ala Asn Ser Thr Gly
165 170 175
Gln His His Leu Pro Arg Ala Thr Cys Ser Met Ser Pro Gly Cys Trp
180 185 190
Gly Gln His His Val Asp Ala Gln Ala Leu Val Gln Pro Asp Val Pro
195 200 205
Lys Ala Asp Ser Gln Gly Pro Gly Arg Glu Pro Glu Gly Pro Met Pro
210 215 220
Ser Gly *
225 226

<210> 1732
<211> 102
<212> PRT
<213> Homo sapiens

<400> 1732
Met Val Ser Lys Phe Leu Leu Ser His Leu Val Leu Ala Val Pro Leu
1 5 10 15
Arg Val Leu Leu Val Leu Trp Ala Leu Cys Val Gly Leu Ser Arg Val
20 25 30
Met Ile Gly Arg His His Val Thr Asp Val Leu Ser Gly Phe Val Ile
35 40 45
Gly Tyr Leu Gln Phe Arg Met Met Glu Lys Val Ser Met Gln Tyr Lys
50 55 60
Thr Cys Arg Met Leu Ile Phe Val Trp Arg Arg Ala Arg Arg Pro Thr
65 70 75 80
His Thr Phe Glu Gly Arg Leu Val Ser Lys Lys Gly Gln Asp Leu Ala
85 90 95
Arg Trp Leu Ser Leu *
100 101

<210> 1733
<211> 139
<212> PRT
<213> Homo sapiens

<400> 1733
Met Lys Phe Thr Thr Leu Leu Phe Leu Ala Ala Val Ala Gly Ala Leu
1 5 10 15
Val Tyr Ala Glu Asp Ala Ser Ser Asp Ser Thr Gly Ala Asp Pro Ala
20 25 30
Gln Glu Ala Gly Thr Ser Lys Pro Asn Glu Glu Ile Ser Gly Pro Ala
35 40 45
Glu Pro Ala Ser Pro Pro Glu Thr Thr Thr Thr Ala Gln Glu Thr Ser
50 55 60
Ala Ala Ala Val Gln Gly Thr Ala Lys Val Thr Ser Ser Arg Gln Glu
65 70 75 80
Leu Asn Pro Leu Lys Ser Ile Val Glu Lys Ser Ile Leu Leu Thr Glu
85 90 95
Gln Ala Leu Ala Lys Ala Gly Lys Gly Met His Gly Gly Val Pro Gly
100 105 110
Gly Lys Gln Phe Ile Glu Asn Gly Ser Glu Phe Ala Gln Lys Leu Leu
115 120 125
Lys Lys Phe Ser Leu Leu Lys Pro Trp Ala *
130 135 138

<210> 1734
<211> 60
<212> PRT
<213> Homo sapiens

<400> 1734
Met Val Arg Ala Ser Phe Leu Cys Cys Val His Arg Thr Leu Gly Pro
1 5 10 15
Trp Asp Leu Ser His Met Glu Leu Gly Gln Leu Leu Gln Asn Ala Pro
20 25 30
Ser Ala His Arg Gly Cys Leu Gly Val Trp Lys Glu Val Val Pro Lys

```
        35                  40                  45
Gln Leu Val Cys Trp Ile Leu Thr Phe Phe Phe  *
    50                  55              59


     <210> 1735
     <211> 73
     <212> PRT
     <213> Homo sapiens


     <400> 1735
Met Cys Ala Cys Ala Val Arg Ala Leu Ser Leu Ala Gly Gly Ala Val
 1               5                  10                  15
Leu Leu Ser Ser Leu Cys Ala Cys Ala Arg Ala Pro Arg Tyr Val Gly
            20                  25                  30
Gly Glu Arg Arg Val Gln Ser Pro Ala Arg Pro Ala Asp Ser Val Ala
        35                  40                  45
Arg Ile Ala Phe Ile Leu Phe Arg Phe Arg Thr Asp Leu Gln Ser Gly
    50                  55                  60
Pro Ser Leu His Leu Gly Ile Cys  *
 65                 70      72


     <210> 1736
     <211> 65
     <212> PRT
     <213> Homo sapiens


     <400> 1736
Met Met Ala Leu Phe Thr Gly Lys Leu Leu Gln Val Val Ser Lys Val
 1               5                  10                  15
Leu Trp Leu Tyr Gln Thr Asn Phe Ser Leu His Thr His Tyr Ser Phe
            20                  25                  30
Asn Arg Gly Gln Ile Phe Lys Arg Lys Thr Val Gln Asn Cys Arg His
        35                  40                  45
Thr Cys Ala Asn Pro Gly Ser Val Glu Arg Leu Ile Trp Glu Phe Gln
    50                  55                  60              64
 *


     <210> 1737
     <211> 47
     <212> PRT
     <213> Homo sapiens


     <400> 1737
Met Ile Gln Val Arg Asn Leu Ile Val Leu Val Cys Phe Leu Val Glu
 1               5                  10                  15
Leu Leu Asn Val Pro Val Leu Phe Leu Tyr Ser Arg Gly Trp Gln Thr
            20                  25                  30
Leu Thr His Gly Leu Thr Gln Leu Lys Thr Ala Phe Phe Leu  *
        35                  40                  45  46
```

<210> 1738
<211> 107
<212> PRT
<213> Homo sapiens

<400> 1738
Met Val Thr Gln Leu Thr Leu Glu Val Leu His Leu Ser Leu Val Val
1 5 10 15
Gly Gln Val Ser Asn Asn Leu Leu Leu His Ile Gly Pro Leu Ala Ser
20 25 30
Glu Gln Met Phe Tyr Ala Val Ala Thr Lys Ile Arg Asp Glu Asn Thr
35 40 45
Tyr Lys Ile Cys Thr Trp Leu Glu Ile Lys Val His His Val Leu Leu
50 55 60
His Ile Gln Gly Thr Leu Thr Cys Ser Tyr Leu Ser His Ser Glu Gln
65 70 75 80
Leu Val Phe Gln Ser Tyr Glu Tyr Val Asp Cys Arg Gly Asn Ala Ser
85 90 95
Val Pro His Gln Leu Thr Pro His Pro Pro *
100 105 106

<210> 1739
<211> 90
<212> PRT
<213> Homo sapiens

<400> 1739
Met Val Leu Pro Pro His Lys Thr Val Gln Leu Pro Arg Leu His Leu
1 5 10 15
Val Trp Leu Trp Val Ser Gln Ala Trp Val Gly Gly Thr Val Leu His
20 25 30
Trp Leu Ala Ser Gln Gln Leu Cys Val Leu Val Pro Ala Ser Leu Thr
35 40 45
Met Ser Trp Asp Leu Glu Ala Arg Leu Gly Tyr Ile Leu Ala Trp Met
50 55 60
Ser Leu Gly Pro Cys Tyr Cys Cys Leu Phe Thr Ile Pro Thr Leu Leu
65 70 75 80
Glu Ile Ser Leu Ile Val Ser Leu Ala *
85 89

<210> 1740
<211> 57
<212> PRT
<213> Homo sapiens

<400> 1740
Met His Cys Val Leu Glu Ile Leu Val Ser Val Leu Gly Leu Thr His
1 5 10 15
His Leu Leu Leu Arg Asp Arg Asp His Tyr Arg Leu Val Arg Leu Met

```
            20                  25                  30
Gly Asp Val Gly Gly Glu Gly Glu Leu Lys Ala Met Trp Arg Val Cys
        35                  40                  45
Leu Ser Val Cys Arg Val Asp Lys  *
    50                  55  56
```

```
     <210> 1741
     <211> 49
     <212> PRT
     <213> Homo sapiens
```

```
     <400> 1741
Met Ile Leu Asn Lys Ala Leu Met Leu Gly Ala Leu Ala Leu Thr Thr
 1               5                  10                  15
Val Met Ser Pro Cys Gly Gly Glu Gly Ile Val Gly Glu Cys Met Ser
            20                  25                  30
Glu Gly Cys Ser Leu Glu Leu Lys Asn Ser Lys Leu Lys Glu Lys Arg
        35                  40                  45              48
 *
```

```
     <210> 1742
     <211> 87
     <212> PRT
     <213> Homo sapiens
```

```
     <400> 1742
Met Ser Phe Val Lys Ile Leu Ile Trp Glu Leu Phe Ile Ala Cys Phe
 1               5                  10                  15
Pro Gln Gly Pro Leu Val His Ser Gly Lys Met Leu Lys His Gly Leu
            20                  25                  30
Asp Trp His Arg Thr Leu Leu Gln Lys His Pro Cys Ile Leu Phe Phe
        35                  40                  45
Ser Phe Leu Lys Trp Asn Leu Ala Leu Ser Pro Trp Met Glu Gly Ser
    50                  55                  60
Gly Ala Ile Ser Ala His Cys Asn Leu Cys Leu Leu Gly Ser Arg Asp
65                  70                  75                  80
Ala Pro Ala Ser Val Ser  *
                    85  86
```

```
     <210> 1743
     <211> 49
     <212> PRT
     <213> Homo sapiens
```

```
     <400> 1743
Met Gly Phe Leu Ser Leu Thr Leu Tyr Leu Leu Thr Ser Leu Asn Lys
 1               5                  10                  15
Met Leu Phe Lys Leu Arg Gly Ala Gln Pro Thr Glu Glu Asp Ile Gly
            20                  25                  30
```

```
Gly Trp Leu Asn Glu Leu Lys Thr Ser Leu Lys Tyr Ile Arg Leu Arg
        35                  40                  45          48
 *
```

<210> 1744
<211> 57
<212> PRT
<213> Homo sapiens

<400> 1744
```
Met Gly Val Ser Glu Leu Leu Leu Leu Leu Lys Met Ile Ala Ser Val
  1               5                  10                  15
Ile Phe Leu Tyr Ser Phe Ile Ser Met Phe Lys Thr Gln Leu Leu Cys
             20                  25                  30
Ser Ser Ser Thr Ser His Gly Ile Leu Glu Ser Arg Ile Lys Cys His
        35                  40                  45
Ala Asp Phe Tyr Leu Phe Cys Gln  *
    50                  55  56
```

<210> 1745
<211> 96
<212> PRT
<213> Homo sapiens

<400> 1745
```
Met Asn Gln Leu Ser Phe Leu Leu Phe Leu Ile Ala Thr Thr Arg Gly
  1               5                  10                  15
Trp Ser Thr Asp Glu Ala Asn Thr Tyr Phe Leu Glu Cys Thr Cys Ser
             20                  25                  30
Trp Ser Pro Ser Leu Pro Lys Ser Cys Pro Glu Ile Lys Asp Gln Cys
        35                  40                  45
Pro Ser Ala Phe Asp Gly Leu Tyr Phe Ile Arg Thr Glu Asn Ala Val
    50                  55                  60
Ile His His Thr Phe Cys Val Met Thr Ser Ala Gly Cys Phe Trp Ile
 65                  70                  75                      80
Leu Lys Val Thr Val His Asn Tyr Asp Leu Thr Thr Asp Thr Pro  *
                 85                  90                  95
```

<210> 1746
<211> 53
<212> PRT
<213> Homo sapiens

<400> 1746
```
Met Val Ile Ser Ala Ala Val Leu Ser Ser Ile Leu Cys Val Phe Leu
  1               5                  10                  15
Ser Lys Leu Val Leu Met Asn Asp Glu Cys Leu Arg Leu Thr Phe Trp
             20                  25                  30
Leu His Cys Asn Ala Lys His Tyr Arg Tyr Ser Met Leu Gly Phe Pro
```

```
        35                  40                  45
Lys Leu Thr Ser Val
    50          53


     <210> 1747
     <211> 49
     <212> PRT
     <213> Homo sapiens


     <400> 1747
Met Asn Phe Glu Ile Leu Ile Gln Arg Ser Leu Leu Phe Tyr Phe Val
  1               5                  10                  15
Leu Ala Leu Asn Phe Pro Val Ala Ser Leu Asp Phe Phe Ser Val Lys
             20                  25                  30
Ile Ile Ser Ala Val Phe Val Glu Gln Lys Phe Trp Asp Phe Val Lys
        35                  40                  45          48
 *


     <210> 1748
     <211> 196
     <212> PRT
     <213> Homo sapiens


     <400> 1748
Met Ala Met Leu Pro Phe Pro Ile Phe Leu Val Leu Leu Leu Arg Gly
  1               5                  10                  15
Leu Val Leu Trp Thr Pro Ala Ser Ser Gly Thr Ile Met Pro Glu Glu
             20                  25                  30
Arg Lys Thr Glu Ile Glu Arg Glu Thr Glu Thr Glu Ser Glu Thr Val
        35                  40                  45
Ile Gly Thr Glu Lys Glu Asn Ala Pro Glu Arg Glu Arg Gly Ser Val
    50                  55                  60
Ile Thr Val Leu His Gln Val Phe Ser Thr Ala Met Lys Asn Asp Thr
 65                  70                  75                  80
Asp Thr Gly Asn Met Gln Lys Glu Val Met Ser Val Thr Glu Gln Val
                 85                  90                  95
Glu Lys Lys Lys Asn Asp Ile Glu Lys Asp Asp Thr Gly Arg Lys Arg
            100                 105                 110
Lys Pro Asp Ile Ser Leu Leu Glu Val Ile Val Asp Val Ala Met Lys
        115                 120                 125
Val Lys Lys Glu Ile Val Thr Gly Asp Thr Asn Thr Lys Asn Leu Lys
    130                 135                 140
Glu Ala Lys Lys Glu Lys Lys Arg Ala Val Ser Leu Pro Leu Asn Arg
145                 150                 155                 160
Arg Ala Pro Lys Leu His Leu Gln Asn Arg His Gly Phe Gly Leu Leu
                165                 170                 175
Cys Ile Leu Val Pro Glu Val Asp Thr Ile Asn Leu Val Ile Phe Leu
            180                 185                 190
Asp Asn Val  *
        195
```

<210> 1749
<211> 46
<212> PRT
<213> Homo sapiens

<400> 1749
Met Leu Val Lys Val Val Tyr Val Met Gly Ala Ile Leu Lys Ile Phe
1               5                   10                  15
Leu Arg Glu Gly Asn Val Ile Asn Gln Arg Ser Gly Met Asp Ile Glu
            20                  25                  30
Lys Tyr Ser Glu His Tyr Leu Ala Gln Gly Val Arg Trp  *
        35                  40                  45

<210> 1750
<211> 82
<212> PRT
<213> Homo sapiens

<400> 1750
Met Glu Leu Val Arg Arg Leu Met Pro Leu Thr Leu Leu Ile Leu Ser
1               5                   10                  15
Cys Leu Ala Glu Leu Thr Met Ala Glu Ala Glu Gly Asn Ala Ser Cys
            20                  25                  30
Thr Val Ser Leu Gly Gly Ala Asn Met Ala Glu Thr His Lys Ala Met
        35                  40                  45
Ile Leu Gln Leu Asn Pro Ser Glu Asn Cys Thr Trp Thr Ile Glu Arg
    50                  55                  60
Pro Glu Asn Lys Ser Ile Arg Ile Ile Phe Cys Tyr Val Gln Leu Gly
65                  70                  75                  80
Ser Glu
    82

<210> 1751
<211> 94
<212> PRT
<213> Homo sapiens

<400> 1751
Met Gly Ser Val Phe Trp His Val Leu Phe Cys Ile Ser Gly Val Cys
1               5                   10                  15
Leu Trp Cys Ala His Arg Met Ala Ala Phe Leu Gln Gln Met Ala Val
            20                  25                  30
Leu Leu Pro Val Asp Cys Glu Arg Pro Ala Ala Val His Trp Leu Ala
        35                  40                  45
Leu Cys Gly Cys Cys Tyr Gly Gln Leu Val Trp Glu Ser Arg Thr Arg
    50                  55                  60
Ser Cys Phe Trp Ser Leu Glu Cys Leu Cys Phe Gly Gly Gln His Phe
65                  70                  75                  80
Gly Ser Val Pro Ser Phe Phe Cys Ser Ser Val Trp Leu  *
                85                  90          93

<210> 1752
<211> 143
<212> PRT
<213> Homo sapiens

<400> 1752
Met Asp Thr Trp Leu Val Cys Trp Ala Ile Phe Ser Leu Leu Lys Ala
1 5 10 15
Gly Leu Thr Glu Pro Glu Val Thr Gln Thr Pro Ser His Gln Val Thr
20 25 30
Gln Met Gly Gln Glu Val Ile Leu Arg Cys Val Pro Ile Ser Asn His
35 40 45
Leu Tyr Phe Tyr Trp Tyr Arg Gln Ile Leu Gly Gln Lys Val Glu Phe
50 55 60
Leu Val Ser Phe Tyr Asn Asn Glu Ile Ser Glu Lys Ser Glu Ile Phe
65 70 75 80
Asp Asp Gln Phe Ser Val Glu Arg Pro Asp Gly Ser Asn Phe Thr Leu
85 90 95
Lys Ile Arg Ser Thr Lys Leu Glu Asp Ser Ala Met Tyr Phe Cys Ala
100 105 110
Ser Ser Glu Arg Gly Ser Gly Ala Asn Val Leu Thr Phe Gly Ala Gly
115 120 125
Ser Arg Leu Thr Val Leu Glu Asp Leu Lys Asn Val Phe Pro Pro
130 135 140 143

<210> 1753
<211> 64
<212> PRT
<213> Homo sapiens

<400> 1753
Met Val Cys Arg Leu Pro Cys Thr Leu Leu Pro Trp Pro Leu Lys His
1 5 10 15
Lys Gln Gly Ala Leu Leu Tyr Ile Cys Pro Ala Ser Leu Pro Ala Phe
20 25 30
Asn Pro Arg Asn Leu Ser Val Tyr Leu Leu Phe Ser Ala Ser Glu Ser
35 40 45
Leu Pro Leu Lys Ser Glu Gln Ala Arg Pro Gly Gly Ser Arg Leu *
50 55 60 63

<210> 1754
<211> 124
<212> PRT
<213> Homo sapiens

<400> 1754
Met Val Leu Gln Thr His Ala Phe Ile Ser Leu Leu Leu Trp Ile Ser
1 5 10 15
Gly Ala Cys Gly Asp Ile Val Met Thr His Ser Pro Asp Ser Leu Ala
20 25 30

Val Ser Leu Gly Glu Thr Ala Thr Ile Asp Cys Arg Ser Ser Gln Ser
35 40 45
Val Leu Tyr His Ala Asn Asn Lys Asn Tyr Leu Thr Trp Tyr Gln Gln
50 55 60
Arg Pro Arg Gln Ser Pro Lys Val Leu Ile Phe Trp Ala Ser Thr Arg
65 70 75 80
Glu Thr Gly Val Pro Asp Arg Phe Thr Gly Ser Gly Ser Gly Thr Asp
85 90 95
Tyr Ser Leu Thr Ile Ser Ser Leu Gln Ala Glu Asp Val Ala Thr Tyr
100 105 110
Tyr Cys Gln Gln Tyr Tyr Asp Ser Pro Ile Thr Phe
115 120 124

<210> 1755
<211> 111
<212> PRT
<213> Homo sapiens

<400> 1755
Met Gln Ala Thr Ser Asn Leu Leu Asn Leu Leu Leu Leu Ser Leu Phe
1 5 10 15
Ala Gly Leu Asn Pro Ser Lys Thr His Ile Asn Pro Lys Glu Gly Trp
20 25 30
Gln Val Tyr Ser Ser Ala Gln Asp Pro Asp Gly Arg Gly Ile Cys Thr
35 40 45
Val Val Ala Pro Glu Gln Asn Leu Cys Ser Arg Asp Ala Lys Ser Arg
50 55 60
Gln Leu Arg Gln Leu Leu Glu Lys Val Gln Asn Met Ser Gln Ser Ile
65 70 75 80
Glu Val Leu Asn Leu Arg Thr Gln Arg Asp Phe Gln Tyr Val Leu Lys
85 90 95
Met Glu Thr Gln Met Lys Gly Leu Lys Ala Lys Phe Arg Gln Ile
100 105 110 111

<210> 1756
<211> 74
<212> PRT
<213> Homo sapiens

<400> 1756
Met Leu Pro Arg Leu Val Leu Ser Ser Trp Pro Gln Ser Ile Phe Leu
1 5 10 15
Pro Arg Phe Trp Asn Tyr Arg Cys Glu Pro Pro Cys Leu Ala Cys Phe
20 25 30
Asp Ile Phe Tyr Ser Val Leu Ile Thr Asn Ser Leu His Met Pro Glu
35 40 45
Tyr Lys Ser Lys Cys Tyr Leu Leu Phe Arg Trp Glu Leu Gln Lys Leu
50 55 60
His Gln Lys Tyr Ala Leu Arg Tyr Ile *
65 70 73

<210> 1757
<211> 50
<212> PRT
<213> Homo sapiens

<400> 1757
Met Glu Asn Val Asn Leu Lys Ala Ser Tyr Leu Gln Phe Ser Lys Leu
1 5 10 15
Met Ala Gly Lys Gly Trp Ala Leu Phe Ile Ala Leu Thr Phe Ser Gln
20 25 30
Arg Leu Leu Pro Cys Leu Ala Ile Ile Glu Ile Ile Asn Val Gly Val
35 40 45
Glu *
49

<210> 1758
<211> 123
<212> PRT
<213> Homo sapiens

<400> 1758
Met Ala Trp Ile Pro Leu Phe Leu Gly Val Leu Ala Tyr Cys Thr Glu
1 5 10 15
Ser Val Ala Ser Tyr Glu Leu Phe Gln Pro Pro Ser Val Ser Val Ser
20 25 30
Pro Gly Gln Thr Ala Thr Phe Thr Cys Ser Gly Asp Asp Leu Gly Asn
35 40 45
Lys Tyr Ile Cys Trp Tyr Leu Gln Lys Pro Gly Gln Pro Pro Val Val
50 55 60
Leu Met Tyr Gln Asp Asn Lys Arg Pro Ser Gly Ile Pro Glu Arg Phe
65 70 75 80
Ser Gly Ser Asn Ser Gly Ser Thr Ala Thr Leu Thr Ile Ser Gly Thr
85 90 95
Gln Ala Thr Asp Glu Ala Leu Tyr Phe Cys Gln Ala Trp Asp Thr Asn
100 105 110
Gly Ala Val Phe Gly Gly Gly Thr Gln Leu Thr
115 120 123

<210> 1759
<211> 75
<212> PRT
<213> Homo sapiens

<400> 1759
Met Arg Trp Arg Thr Ile Leu Leu Gln Tyr Cys Phe Leu Leu Ile Thr
1 5 10 15
Cys Leu Leu Thr Ala Leu Glu Ala Val Pro Ile Asp Ile Asp Lys Thr
20 25 30
Lys Val Gln Asn Ile His Pro Val Glu Ser Ala Lys Ile Glu Pro Pro
35 40 45
Asp Thr Gly Leu Tyr Tyr Asp Glu Ile Val Leu Glu Glu Leu Gly Gly
50 55 60

Pro Cys Leu Tyr Leu Glu Gly Asn Pro Thr *
65 70 74

<210> 1760
<211> 122
<212> PRT
<213> Homo sapiens

<400> 1760
Met Arg Leu Pro Asp Val Gln Leu Trp Leu Val Leu Leu Trp Ala Leu
1 5 10 15
Val Arg Ala Gln Gly Thr Gly Ser Val Cys Pro Ser Cys Gly Gly Ser
20 25 30
Lys Leu Ala Pro Gln Ala Glu Arg Ala Leu Val Leu Glu Leu Ala Lys
35 40 45
Gln Gln Ile Leu Asp Gly Leu His Leu Thr Ser Arg Pro Arg Ile Thr
50 55 60
His Pro Pro Pro Gln Ala Ala Leu Thr Arg Ala Leu Arg Arg Leu Gln
65 70 75 80
Pro Gly Ser Val Ala Pro Gly Asn Gly Glu Glu Val Ile Ser Phe Ala
85 90 95
Thr Val Thr Asp Ser Thr Ser Ala Tyr Ser Ser Leu Leu Thr Phe His
100 105 110
Leu Ser Thr Pro Arg Ser His His Leu Tyr
115 120 122

<210> 1761
<211> 123
<212> PRT
<213> Homo sapiens

<400> 1761
Met Arg Val Arg Ile Gly Leu Thr Leu Leu Leu Cys Ala Val Leu Leu
1 5 10 15
Ser Leu Ala Ser Ala Ser Ser Asp Glu Glu Gly Ser Gln Asp Glu Ser
20 25 30
Leu Asp Ser Lys Thr Thr Leu Thr Ser Asp Glu Ser Val Lys Asp His
35 40 45
Thr Thr Ala Gly Arg Val Val Ala Gly Gln Ile Phe Leu Asp Ser Glu
50 55 60
Glu Ser Glu Leu Glu Ser Ser Ile Gln Glu Glu Glu Asp Ser Leu Lys
65 70 75 80
Ser Gln Glu Gly Glu Ser Val Thr Glu Asp Ile Ser Phe Leu Glu Ser
85 90 95
Pro Asn Pro Glu Asn Lys Asp Tyr Glu Glu Pro Lys Lys Val Arg Lys
100 105 110
Pro Gly Ser Leu Asp Ile Phe Leu Ala Phe *
115 120 122

<210> 1762
<211> 145

<212> PRT
<213> Homo sapiens

<221> misc_feature
<222> (1)...(145)
<223> Xaa = any amino acid or nothing

<400> 1762
Met Ala Leu Ala Ala Leu Met Ile Ala Leu Gly Ser Leu Gly Leu His
1 5 10 15
Thr Trp Gln Ala Gln Ala Val Pro Thr Ile Leu Pro Leu Gly Leu Ala
20 25 30
Pro Asp Thr Phe Asp Asp Thr Tyr Val Gly Cys Ala Glu Glu Met Glu
35 40 45
Glu Lys Ala Ala Pro Leu Leu Lys Glu Glu Met Ala His His Ala Leu
50 55 60
Leu Arg Glu Ser Trp Glu Ala Ala Gln Glu Thr Trp Glu Asp Lys Arg
65 70 75 80
Arg Gly Leu Thr Leu Pro Pro Gly Phe Lys Ala Gln Asn Gly Ile Ala
85 90 95
Ile Met Val Tyr Thr Asn Ser Ser Asn Thr Leu Tyr Trp Glu Leu Asn
100 105 110
Xaa Ala Val Arg Thr Gly Gly Gly Ser Arg Glu Leu Tyr Met Arg His
115 120 125
Phe Pro Phe Lys Ala Leu His Phe Tyr Leu Ile Arg Ala Leu Gln Leu
130 135 140
Leu
145

<210> 1763
<211> 257
<212> PRT
<213> Homo sapiens

<400> 1763
Met Lys Arg Glu Arg Gly Ala Leu Ser Arg Ala Ser Arg Ala Leu Arg
1 5 10 15
Leu Ala Pro Phe Val Tyr Leu Leu Leu Ile Gln Thr Asp Pro Leu Glu
20 25 30
Gly Val Asn Ile Thr Ser Pro Val Arg Leu Ile His Gly Thr Val Gly
35 40 45
Lys Ser Ala Leu Leu Ser Val Gln Tyr Ser Ser Thr Ser Ser Asp Arg
50 55 60
Pro Val Val Lys Trp Gln Leu Lys Arg Asp Lys Pro Val Thr Val Val
65 70 75 80
Gln Ser Ile Gly Thr Glu Val Ile Gly Thr Leu Arg Pro Asp Tyr Arg
85 90 95
Asp Arg Ile Arg Leu Phe Glu Asn Gly Ser Leu Leu Leu Ser Asp Leu
100 105 110
Gln Leu Ala Asp Glu Gly Thr Tyr Glu Val Glu Ile Ser Ile Thr Asp
115 120 125
Asp Thr Phe Thr Gly Glu Lys Thr Ile Asn Leu Thr Val Asp Val Pro
130 135 140
Ile Ser Arg Pro Gln Val Leu Gly Ala Ser Thr Thr Val Leu Glu Leu
145 150 155 160

```
Ser Glu Ala Phe Thr Leu Asn Cys Ser His Glu Asn Gly Thr Lys Pro
                165                 170                 175
Ser Tyr Thr Trp Leu Lys Asp Gly Lys Pro Leu Leu Asn Asp Ser Arg
            180                 185                 190
Met Leu Leu Ser Pro Asp Gln Lys Val Leu Thr Ile Thr Arg Val Leu
        195                 200                 205
Met Glu Asp Asp Asp Leu Tyr Ser Cys Val Val Glu Asn Pro Ile Asn
    210                 215                 220
Gln Gly Arg Thr Leu Pro Cys Lys Ile Thr Glu Tyr Arg Lys Ser Ser
225                 230                 235                 240
Leu Ser Ser Ile Trp Leu Gln Glu Ala Phe Ser Ser Leu Gly Pro Trp
                245                 250                 255 256
 *
```

```
<210> 1764
<211> 166
<212> PRT
<213> Homo sapiens

<221> misc_feature
<222> (1)...(166)
<223> Xaa = any amino acid or nothing


<400> 1764
Met Ala Leu Lys Val Leu Leu Glu Gln Glu Lys Thr Phe Phe Thr Leu
 1               5                  10                  15
Leu Val Leu Leu Gly Tyr Leu Ser Cys Lys Val Thr Cys Glu Ser Gly
            20                  25                  30
Asp Cys Arg Gln Gln Glu Phe Arg Asp Arg Ser Gly Asn Cys Val Pro
        35                  40                  45
Cys Asn Gln Cys Gly Pro Gly Met Glu Leu Ser Lys Glu Cys Gly Phe
    50                  55                  60
Gly Tyr Gly Glu Asp Ala Gln Cys Val Thr Cys Arg Leu His Arg Phe
65                  70                  75                  80
Lys Glu Asp Trp Gly Phe Gln Lys Cys Lys Pro Cys Leu Asp Cys Ala
                85                  90                  95
Val Val Asn Arg Phe Gln Lys Ala Asn Cys Ser Ala Thr Ser Asp Ala
            100                 105                 110
Ile Cys Gly Asp Cys Leu Pro Gly Phe Tyr Arg Lys Thr Lys Leu Val
        115                 120                 125
Gly Phe Gln Asp Met Glu Trp Trp Xaa Ala Leu Val Gly Arg Thr Pro
    130                 135                 140
Phe Leu Pro Ser Leu Tyr Gly Asn Pro Ala Leu Gly Cys Gln Pro Arg
145                 150                 155                 160
Val Gln Thr Phe Gly Glu
                165 166


<210> 1765
<211> 90
<212> PRT
<213> Homo sapiens
```

<400> 1765
Met Ser Cys Ser Cys Pro Pro Cys Phe Phe Thr Leu Phe Leu His Ser
1               5                   10                  15
Ile Cys Gln Asp Ile Ser Trp Phe His Pro Gln Thr Pro Thr Leu Asp
            20                  25                  30
Ser Leu Leu Asn Trp Ile Asp Asp Leu Ile Phe Tyr Gly Thr Leu Tyr
        35                  40                  45
Asn Phe Phe Pro Glu Glu Thr Pro Leu Phe Thr Phe Leu Leu Thr Leu
    50                  55                  60
Tyr Leu Ser Leu Leu Leu Leu Trp Leu Pro Gly Met Ala Ala Leu Pro
65                  70                  75                  80
Leu Ala Val Met Pro Asn Tyr Leu Tyr Lys
                85                  90

<210> 1766
<211> 57
<212> PRT
<213> Homo sapiens

<400> 1766
Met Pro Ala Leu Arg Pro Ala Leu Leu Trp Ala Leu Leu Ser Leu Trp
1               5                   10                  15
Leu Cys Cys Ala Thr Pro Ala Pro Ala Leu Gln Cys Pro Glu Gly Tyr
            20                  25                  30
Glu Pro Ser Pro Leu Asp Arg Lys Cys Ala Pro Tyr Pro Asn Val Arg
        35                  40                  45
Arg Ser Cys Pro Cys Pro Glu Gly Phe
    50                  55      57

<210> 1767
<211> 63
<212> PRT
<213> Homo sapiens

<400> 1767
Met Val Phe Leu Tyr Gly Phe Val Phe Ile Lys Lys Ala Gln Leu Ile
1               5                   10                  15
Val Val Leu Leu Phe Thr Asp Val Ala Gln Arg Thr Ala Ala Gly Arg
            20                  25                  30
Pro Pro Thr Pro Val Leu Gly Pro Pro Ser Pro Glu Cys Cys Leu Leu
        35                  40                  45
Phe Met Glu Gly Glu Gln Trp Ile Leu Gly Thr Thr Gly Gln Ala
    50                  55                  60          63

<210> 1768
<211> 174
<212> PRT
<213> Homo sapiens

<400> 1768

Met Pro Ser Gly Cys Arg Cys Leu His Leu Val Cys Leu Leu Cys Ile
1 5 10 15
Leu Gly Ala Pro Gly Gln Pro Val Arg Ala Asp Asp Cys Ser Ser His
20 25 30
Cys Asp Leu Ala His Gly Cys Cys Ala Pro Asp Gly Ser Cys Arg Cys
35 40 45
Asp Pro Gly Trp Glu Gly Leu His Cys Glu Arg Cys Val Arg Met Pro
50 55 60
Gly Cys Gln His Gly Thr Cys His Gln Pro Trp Gln Cys Ile Cys His
65 70 75 80
Ser Gly Trp Ala Gly Lys Phe Cys Asp Lys Asp Glu His Ile Cys Thr
85 90 95
Thr Gln Ser Pro Cys Gln Asn Gly Gly Gln Cys Met Tyr Asp Gly Gly
100 105 110
Gly Glu Tyr His Cys Val Cys Leu Pro Gly Phe His Gly Arg Asp Cys
115 120 125
Glu Arg Lys Ala Gly Pro Cys Glu Gln Ala Gly Ser Pro Cys Arg Asn
130 135 140
Gly Gly Gln Cys Gln Asp Asp Gln Gly Phe Ala Leu Asn Phe Thr Cys
145 150 155 160
Arg Cys Leu Val Gly Phe Val Gly Ala Arg Cys Asp Val *
165 170 173

<210> 1769
<211> 78
<212> PRT
<213> Homo sapiens

<400> 1769
Met Leu Cys Leu Cys Arg Phe Ala Cys Ser Arg Arg Phe Thr Ala Met
1 5 10 15
Gly Leu Phe Cys Leu Ala Ser Leu Thr Leu His His Ile Phe Lys Val
20 25 30
His Pro Ser Cys Ser Val Ser Val Pro Pro Gly Phe Ser Leu Leu Ser
35 40 45
Ser Ala Arg Cys Met Asp Arg Pro Arg Cys Ala His Leu Phe Ala Leu
50 55 60
Met Gly Pro Cys Leu Gly Leu Ser Thr Phe Gly Arg Leu *
65 70 75 77

<210> 1770
<211> 149
<212> PRT
<213> Homo sapiens

<400> 1770
Met Leu Val Thr Leu Gly Leu Leu Thr Ser Phe Phe Ser Phe Leu Tyr
1 5 10 15
Met Val Ala Pro Ser Ile Arg Lys Phe Phe Ala Gly Gly Val Cys Arg
20 25 30
Thr Asn Val Gln Leu Pro Gly Lys Val Val Val Ile Thr Gly Ala Asn
35 40 45
Thr Gly Ile Gly Lys Glu Thr Ala Arg Glu Leu Ala Ser Arg Gly Ala

50 55 60
Arg Val Tyr Ile Ala Cys Arg Asp Val Leu Lys Gly Glu Ser Ala Ala
65 70 75 80
Ser Glu Ile Arg Val Asp Thr Lys Asn Ser Gln Val Leu Val Arg Lys
85 90 95
Leu Asp Leu Ser Asp Thr Lys Ser Ile Arg Ala Phe Ala Glu Gly Phe
100 105 110
Leu Ala Glu Glu Lys Gln Leu His Ile Leu Ile Asn Asn Ala Gly Val
115 120 125
Met Met Cys Pro Tyr Ser Lys Thr Ala Asp Gly Phe Glu Thr His Leu
130 135 140
Gly Val Asn His Leu
145 149

<210> 1771
<211> 76
<212> PRT
<213> Homo sapiens

<400> 1771
Met Met Thr Leu Leu Arg Arg Gln Glu Arg Phe Pro Gly Ile Thr Phe
1 5 10 15
Trp Leu Leu Ile Gln Leu Leu Gln Gln Ile Leu Ile Ser Tyr His Gln
20 25 30
Gly Ser Leu Thr Phe Met Glu Asn Gly Asn Cys Leu Leu Gln Leu Phe
35 40 45
Gln Leu Gly Lys Leu Leu Val Gln Ala Ser His Leu His Gly Gln Leu
50 55 60
Leu Val Phe Val Gln Lys Ile Ile Ile Ser Met *
65 70 75

<210> 1772
<211> 128
<212> PRT
<213> Homo sapiens

<400> 1772
Met Gly Ser Thr Lys His Trp Gly Glu Trp Leu Leu Asn Leu Lys Val
1 5 10 15
Ala Pro Ala Gly Val Phe Gly Val Ala Phe Leu Ala Arg Val Ala Leu
20 25 30
Val Phe Tyr Gly Val Phe Gln Asp Arg Thr Leu His Val Arg Tyr Thr
35 40 45
Asp Ile Asp Tyr Gln Val Phe Thr Asp Ala Ala Arg Phe Val Thr Glu
50 55 60
Gly Arg Ser Pro Tyr Leu Arg Ala Thr Tyr Arg Tyr Thr Pro Leu Leu
65 70 75 80
Gly Trp Leu Leu Thr Pro Asn Ile Tyr Leu Ser Glu Leu Phe Gly Lys
85 90 95
Phe Leu Phe Ile Ser Cys Asp Leu Leu Thr Ala Phe Leu Leu Tyr Arg
100 105 110
Leu Leu Leu Leu Lys Gly Leu Gly Arg Arg Gln Ala Cys Gly Tyr Cys
115 120 125 128

<210> 1773
<211> 614
<212> PRT
<213> Homo sapiens

<400> 1773
Met Gly Ala Leu Arg Pro Thr Leu Leu Pro Pro Ser Leu Pro Leu Leu
1 5 10 15
Leu Leu Leu Met Leu Gly Met Gly Cys Trp Ala Arg Glu Val Leu Val
20 25 30
Pro Glu Gly Pro Leu Tyr Arg Val Ala Gly Thr Ala Val Ser Ile Ser
35 40 45
Cys Asn Val Thr Gly Tyr Glu Gly Pro Ala Gln Gln Asn Phe Glu Trp
50 55 60
Phe Leu Tyr Arg Pro Glu Ala Pro Asp Thr Ala Leu Gly Ile Val Ser
65 70 75 80
Thr Lys Asp Thr Gln Phe Ser Tyr Ala Val Phe Lys Ser Arg Val Val
85 90 95
Ala Gly Glu Val Gln Val Gln Arg Leu Gln Gly Asp Ala Val Val Leu
100 105 110
Lys Ile Ala Arg Leu Gln Ala Gln Asp Ala Gly Ile Tyr Glu Cys His
115 120 125
Thr Pro Ser Thr Asp Thr Arg Tyr Leu Gly Ser Tyr Ser Gly Lys Val
130 135 140
Glu Leu Arg Val Leu Pro Asp Val Leu Gln Val Ser Ala Ala Pro Pro
145 150 155 160
Gly Pro Arg Gly Arg Gln Ala Pro Thr Ser Pro Pro Arg Met Thr Val
165 170 175
His Glu Gly Gln Glu Leu Ala Leu Gly Cys Leu Ala Arg Thr Ser Thr
180 185 190
Gln Lys His Thr His Leu Ala Val Ser Phe Gly Arg Ser Val Pro Glu
195 200 205
Ala Pro Val Gly Arg Ser Thr Leu Gln Glu Val Val Gly Ile Arg Ser
210 215 220
Asp Leu Ala Val Glu Ala Gly Ala Pro Tyr Ala Glu Arg Leu Ala Ala
225 230 235 240
Gly Glu Leu Arg Leu Gly Lys Glu Gly Thr Asp Arg Tyr Arg Met Val
245 250 255
Val Gly Gly Ala Gln Ala Gly Asp Ala Gly Thr Tyr His Cys Thr Ala
260 265 270
Ala Glu Trp Ile Gln Asp Pro Asp Gly Ser Trp Ala Gln Ile Ala Glu
275 280 285
Lys Arg Ala Val Leu Ala His Val Asp Val Gln Thr Leu Ser Ser Gln
290 295 300
Leu Ala Val Thr Val Gly Pro Gly Glu Arg Arg Ile Gly Pro Gly Glu
305 310 315 320
Pro Leu Glu Leu Leu Cys Asn Val Ser Gly Ala Leu Pro Pro Ala Gly
325 330 335
Arg His Ala Ala Tyr Ser Val Gly Trp Glu Met Ala Pro Ala Gly Ala
340 345 350
Pro Gly Pro Gly Arg Leu Val Ala Gln Leu Asp Thr Glu Gly Val Gly
355 360 365
Ser Leu Gly Pro Gly Tyr Glu Gly Arg His Ile Ala Met Glu Lys Val

```
    370                 375                 380
Ala Ser Arg Thr Tyr Arg Leu Arg Leu Glu Ala Ala Arg Pro Gly Asp
385                 390                 395                 400
Ala Gly Thr Tyr Arg Cys Leu Ala Lys Ala Tyr Val Arg Gly Ser Gly
                405                 410                 415
Thr Arg Leu Arg Glu Ala Ala Ser Ala Arg Ser Arg Pro Leu Pro Val
            420                 425                 430
His Val Arg Glu Glu Gly Val Val Leu Glu Ala Val Ala Trp Leu Ala
        435                 440                 445
Gly Gly Thr Val Tyr Arg Gly Glu Thr Ala Ser Leu Leu Cys Asn Ile
    450                 455                 460
Ser Val Arg Gly Gly Pro Pro Gly Leu Arg Leu Ala Ala Ser Trp Trp
465                 470                 475                 480
Val Glu Arg Pro Glu Asp Gly Glu Leu Ser Ser Val Pro Ala Gln Leu
                485                 490                 495
Val Gly Gly Val Gly Gln Asp Gly Val Ala Glu Leu Gly Val Arg Pro
            500                 505                 510
Gly Gly Gly Pro Val Ser Val Glu Leu Val Gly Pro Arg Ser His Arg
        515                 520                 525
Leu Arg Leu His Ser Leu Gly Pro Glu Asp Glu Gly Val Tyr His Cys
    530                 535                 540
Ala Pro Ser Ala Trp Val Gln His Ala Asp Tyr Ser Trp Tyr Gln Ala
545                 550                 555                 560
Gly Ser Ala Arg Ser Gly Pro Val Thr Val Tyr Pro Tyr Met His Ala
                565                 570                 575
Leu Asp Thr Leu Phe Val Pro Leu Leu Val Gly Thr Gly Val Ala Leu
            580                 585                 590
Val Thr Gly Ala Thr Val Leu Gly Thr Ile Thr Cys Cys Phe Met Lys
        595                 600                 605
Arg Leu Arg Lys Arg  *
    610         613


     <210> 1774
     <211> 156
     <212> PRT
     <213> Homo sapiens


     <400> 1774
Met Glu Ala Leu Thr Leu Trp Leu Leu Pro Trp Ile Cys Gln Cys Val
 1               5                  10                  15
Ser Val Arg Ala Asp Ser Ile Ile His Ile Gly Ala Ile Phe Glu Glu
            20                  25                  30
Asn Ala Ala Lys Asp Asp Arg Val Phe Gln Leu Ala Val Ser Asp Leu
        35                  40                  45
Ser Leu Asn Asp Asp Ile Leu Gln Ser Glu Lys Ile Thr Tyr Ser Ile
    50                  55                  60
Lys Val Ile Glu Ala Asn Asn Pro Phe Gln Ala Val Gln Glu Ala Cys
65                  70                  75                  80
Asp Leu Met Thr Gln Gly Ile Leu Ala Leu Val Thr Ser Thr Gly Cys
                85                  90                  95
Ala Ser Ala Asn Ala Leu Gln Ser Leu Thr Asp Ala Met His Ile Pro
            100                 105                 110
His Leu Phe Val Gln Arg Asn Pro Gly Gly Ser Pro Arg Thr Ala Cys
        115                 120                 125
His Leu Asn Pro Ser Pro Asp Gly Glu Ala Tyr Thr Leu Ala Ser Arg
    130                 135                 140
```

Pro Pro Val Arg Leu Asn Asp Val Met Leu Arg Leu
145 150 155 156

<210> 1775
<211> 896
<212> PRT
<213> Homo sapiens

<400> 1775
Met Gln Lys Ala Ser Val Leu Leu Phe Leu Ala Trp Val Cys Phe Leu
1 5 10 15
Phe Tyr Ala Gly Ile Ala Leu Phe Thr Ser Gly Phe Leu Leu Thr Arg
20 25 30
Leu Glu Leu Thr Asn His Ser Ser Cys Gln Glu Pro Pro Gly Pro Gly
35 40 45
Ser Leu Pro Trp Gly Ser Gln Gly Lys Pro Gly Ala Cys Trp Met Ala
50 55 60
Ser Arg Phe Ser Arg Val Val Leu Val Leu Ile Asp Ala Leu Arg Phe
65 70 75 80
Asp Phe Ala Gln Pro Gln His Ser His Val Pro Arg Glu Pro Pro Val
85 90 95
Ser Leu Pro Phe Leu Gly Lys Leu Ser Ser Leu Gln Arg Ile Leu Glu
100 105 110
Ile Gln Pro His His Ala Arg Leu Tyr Arg Ser Gln Val Asp Pro Pro
115 120 125
Thr Thr Thr Met Gln Arg Leu Lys Ala Leu Thr Thr Gly Ser Leu Pro
130 135 140
Thr Phe Ile Asp Ala Gly Ser Asn Phe Ala Ser His Ala Ile Val Glu
145 150 155 160
Asp Asn Leu Ile Lys Gln Leu Thr Ser Ala Gly Arg Arg Val Val Phe
165 170 175
Met Gly Asp Asp Thr Trp Lys Asp Leu Phe Pro Gly Ala Phe Ser Lys
180 185 190
Ala Phe Phe Phe Pro Ser Phe Asn Val Arg Asp Leu Asp Thr Val Asp
195 200 205
Asn Gly Ile Leu Glu His Leu Tyr Pro Thr Met Asp Ser Gly Glu Trp
210 215 220
Asp Val Leu Ile Ala His Phe Leu Gly Val Asp His Cys Gly His Lys
225 230 235 240
His Gly Pro His His Pro Glu Met Ala Lys Lys Leu Ser Gln Met Asp
245 250 255
Gln Val Ile Gln Gly Leu Val Glu Arg Leu Glu Asn Asp Thr Leu Leu
260 265 270
Val Val Ala Gly Asp His Gly Met Thr Thr Asn Gly Asp His Gly Gly
275 280 285
Asp Ser Glu Leu Glu Val Ser Ala Ala Leu Phe Leu Tyr Ser Pro Thr
290 295 300
Ala Val Phe Pro Ser Thr Pro Pro Glu Glu Pro Glu Val Ile Pro Gln
305 310 315 320
Val Ser Leu Val Pro Thr Leu Ala Leu Leu Leu Gly Leu Pro Ile Pro
325 330 335
Phe Gly Asn Ile Gly Glu Val Met Ala Glu Leu Phe Ser Gly Gly Glu
340 345 350
Asp Ser Gln Pro His Ser Ser Ala Leu Ala Gln Ala Ser Ala Leu His
355 360 365
Leu Asn Ala Gln Gln Val Ser Arg Phe Phe His Thr Tyr Ser Ala Ala

370 375 380
Thr Gln Asp Leu Gln Ala Lys Glu Leu His Gln Leu Gln Asn Leu Phe
385 390 395 400
Ser Lys Ala Ser Ala Asp Tyr Gln Trp Leu Leu Gln Ser Pro Lys Gly
405 410 415
Ala Glu Ala Thr Leu Pro Thr Val Ile Ala Glu Leu Gln Gln Phe Leu
420 425 430
Arg Gly Ala Arg Ala Met Cys Ile Glu Ser Trp Ala Arg Phe Ser Leu
435 440 445
Val Arg Met Ala Gly Gly Thr Ala Leu Leu Ala Ala Ser Cys Phe Ile
450 455 460
Cys Leu Leu Ala Ser Gln Trp Ala Ile Ser Pro Gly Phe Pro Phe Cys
465 470 475 480
Pro Leu Leu Leu Thr Pro Val Ala Trp Gly Leu Val Gly Ala Ile Ala
485 490 495
Tyr Ala Gly Leu Leu Gly Thr Ile Glu Leu Lys Leu Asp Leu Val Leu
500 505 510
Leu Gly Ala Val Ala Ala Val Ser Ser Phe Leu Pro Phe Leu Trp Lys
515 520 525
Ala Trp Ala Gly Trp Gly Ser Lys Arg Pro Leu Ala Thr Leu Phe Pro
530 535 540
Ile Pro Gly Pro Val Leu Leu Leu Leu Leu Phe Arg Leu Ala Val Phe
545 550 555 560
Phe Ser Asp Ser Phe Val Val Ala Glu Ala Arg Ala Thr Pro Phe Leu
565 570 575
Leu Gly Ser Phe Ile Leu Leu Leu Val Val Gln Leu His Trp Glu Gly
580 585 590
Gln Leu Leu Pro Pro Lys Leu Leu Thr Met Pro Arg Leu Gly Thr Ser
595 600 605
Ala Thr Thr Asn Pro Pro Arg His Asn Gly Ala Tyr Ala Leu Arg Leu
610 615 620
Gly Ile Gly Leu Leu Leu Cys Thr Arg Leu Ala Gly Leu Phe His Arg
625 630 635 640
Cys Pro Glu Glu Thr Pro Val Cys His Ser Ser Pro Trp Leu Ser Pro
645 650 655
Leu Ala Ser Met Val Gly Gly Arg Ala Lys Asn Leu Trp Tyr Gly Ala
660 665 670
Cys Val Ala Ala Leu Val Ala Leu Leu Ala Ala Val Arg Leu Trp Leu
675 680 685
Arg Arg Tyr Gly Asn Leu Lys Ser Pro Glu Pro Pro Met Leu Phe Val
690 695 700
Arg Trp Gly Leu Pro Leu Met Ala Leu Gly Thr Ala Ala Tyr Trp Ala
705 710 715 720
Leu Ala Ser Gly Ala Asp Glu Ala Pro Pro Arg Leu Arg Val Leu Val
725 730 735
Ser Gly Ala Ser Met Val Leu Pro Arg Ala Val Ala Gly Leu Ala Ala
740 745 750
Ser Gly Leu Ala Leu Leu Leu Trp Lys Pro Val Thr Val Leu Val Lys
755 760 765
Ala Gly Ala Gly Ala Pro Arg Thr Arg Thr Val Leu Thr Pro Phe Ser
770 775 780
Gly Pro Pro Thr Ser Gln Ala Asp Leu Asp Tyr Val Val Pro Gln Ile
785 790 795 800
Tyr Arg His Met Gln Glu Glu Phe Arg Gly Arg Leu Glu Arg Thr Lys
805 810 815
Ser Gln Gly Pro Leu Thr Val Ala Ala Tyr Gln Leu Gly Ser Val Tyr
820 825 830
Ser Ala Ala Met Val Thr Ala Leu Thr Leu Leu Ala Phe Pro Leu Leu
835 840 845

```
Leu Leu His Ala Glu Arg Ile Ser Leu Val Phe Leu Leu Leu Phe Leu
    850                 855                 860
Gln Ser Phe Leu Leu Leu His Leu Leu Ala Ala Gly Ile Pro Val Thr
865                 870                 875                 880
Thr Pro Gly Lys Tyr Leu Ser Ser Asp Ser Leu Lys Asp Asn Ser Asp
                885                 890                 895 896
```

```
<210> 1776
<211> 178
<212> PRT
<213> Homo sapiens
```

```
<400> 1776
Met Trp Ala Cys Trp Cys Val Leu Gly Thr Pro Gly Val Ala Met Val
  1               5                  10                  15
Leu Leu His Thr Thr Ile Ser Phe Cys Val Ala Gln Phe Arg Ser Gln
             20                  25                  30
Leu Leu Thr Trp Leu Cys Ser Leu Leu Leu Leu Ser Thr Leu Arg Leu
         35                  40                  45
Gln Gly Val Glu Glu Val Lys Arg Arg Trp Tyr Lys Thr Glu Asn Glu
     50                  55                  60
Tyr Tyr Leu Leu Gln Phe Thr Leu Thr Val Arg Cys Leu Tyr Tyr Thr
 65                  70                  75                  80
Ser Phe Ser Leu Glu Leu Cys Trp Gln Gln Leu Pro Ala Ala Ser Thr
                 85                  90                  95
Ser Tyr Ser Phe Pro Trp Met Leu Ala Tyr Val Phe Tyr Tyr Pro Val
            100                 105                 110
Leu His Asn Gly Pro Ile Leu Ser Phe Ser Glu Phe Ile Lys Gln Arg
        115                 120                 125
Ser Gln Trp Ser Asn Arg Glu Phe Gly Met Glu Val Glu Ser Lys Gly
    130                 135                 140
Pro Gly Ala His Pro Pro Gly Phe Glu Ser Leu Leu Cys Phe Gly Leu
145                 150                 155                 160
Arg Val Leu Ala Glu Leu Leu Thr Leu Leu Met Pro Gln Ser Ser Tyr
                165                 170                 175
Gln  *
177
```

```
<210> 1777
<211> 59
<212> PRT
<213> Homo sapiens
```

```
<400> 1777
Met Pro Thr Tyr Trp Leu Ala Asn Leu Arg Pro Gly Leu Gln Pro Phe
  1               5                  10                  15
Leu Leu His Phe Leu Leu Glu Trp Leu Ala Val Phe Cys Cys Lys Ile
             20                  25                  30
Met Val Leu Ala Ala Ala Gly Leu Leu Pro Thr Leu His Met Ala Ser
         35                  40                  45
Phe Phe Ser Asn Ala Leu Tyr Asn Cys Phe Tyr
```

50 55 59

<210> 1778
<211> 137
<212> PRT
<213> Homo sapiens

<400> 1778
Met Val Ala Pro Gly Leu Val Leu Gly Leu Val Leu Pro Leu Ile Leu
1 5 10 15
Trp Ala Asp Arg Ser Ala Gly Ile Gly Phe Arg Phe Ala Ser Tyr Ile
20 25 30
Asn Asn Asp Met Val Leu Gln Lys Glu Pro Ala Gly Ala Val Ile Trp
35 40 45
Gly Phe Gly Thr Pro Gly Ala Thr Val Thr Val Thr Leu Arg Gln Gly
50 55 60
Gln Glu Thr Ile Met Lys Lys Val Thr Ser Val Lys Ala His Ser Asp
65 70 75 80
Thr Trp Met Val Val Leu Asp Pro Met Lys Pro Gly Gly Pro Phe Glu
85 90 95
Val Met Ala Gln Gln Thr Leu Glu Lys Ile Asn Phe Thr Leu Arg Val
100 105 110
His Asp Val Leu Phe Gly Asp Val Trp Leu Cys Ser Gly Gln Ser Asn
115 120 125
Met Gln Met Thr Val Leu Gln Ile Phe
130 135 137

<210> 1779
<211> 65
<212> PRT
<213> Homo sapiens

<400> 1779
Met Lys Val Phe Phe Leu Asp Glu Ser Trp Pro Gln Trp Arg Phe Ala
1 5 10 15
Ala Gly Leu Leu Ala Leu Ser Phe Gly Gly Pro Ala Trp Lys Phe Leu
20 25 30
Ser Val Gln Arg Val Ile Pro Trp Leu Trp Ala Ala Lys Glu Lys Pro
35 40 45
Leu Gly Pro Leu Ala Thr Pro Pro Arg Leu Asn Pro Lys Val Gly Val
50 55 60 64
*

<210> 1780
<211> 53
<212> PRT
<213> Homo sapiens

<400> 1780

Met Phe His Cys Tyr Trp Phe Arg Cys Leu Ser Pro Gln Thr Leu Leu
1 5 10 15
Cys Lys Cys Phe Ser Lys Gly Arg Thr Asp Trp Asn Cys Gly Ser Ala
20 25 30
Arg Ser His Ser Phe Gln Ser His Phe Phe Ser Ala Ala Leu Ser Ser
35 40 45
Cys Gly Thr Leu *
50 52

<210> 1781
<211> 109
<212> PRT
<213> Homo sapiens

<400> 1781
Met Met His Asn Ile Ile Val Lys Glu Leu Ile Val Thr Phe Phe Leu
1 5 10 15
Gly Ile Thr Val Val Gln Met Leu Ile Ser Val Thr Gly Leu Lys Gly
20 25 30
Val Glu Ala Gln Asn Gly Ser Glu Ser Glu Val Phe Val Gly Lys Tyr
35 40 45
Glu Thr Leu Val Phe Tyr Trp Pro Ser Leu Leu Cys Leu Ala Phe Leu
50 55 60
Leu Gly Arg Phe Leu His Met Phe Val Lys Ala Leu Arg Val His Leu
65 70 75 80
Gly Trp Glu Leu Gln Val Glu Glu Lys Ser Val Leu Glu Val His Gln
85 90 95
Gly Glu His Val Lys Gln Leu Leu Arg Ile Pro Arg Pro
100 105 109

<210> 1782
<211> 58
<212> PRT
<213> Homo sapiens

<400> 1782
Met Ala Ser Thr Trp Ser Leu Glu Arg Val Gly Thr Cys Leu Pro Cys
1 5 10 15
Gly Phe Gly Thr Trp Gln Ser Thr Ala Arg Trp Pro Ser Cys Arg Ser
20 25 30
Thr Ser Met Val Trp Leu Val Trp Pro Ser Leu Leu Ala Pro Ser Thr
35 40 45
Leu Ser Leu Trp Ala Thr Ser Met Thr *
50 55 57

<210> 1783
<211> 102
<212> PRT
<213> Homo sapiens

<400> 1783
Met Leu Ile Pro His Gln Leu Pro Leu Cys Ser Pro Trp Leu Val Gln
1 5 10 15
Ala Met Leu Thr Ile Glu Val Pro Trp Leu Leu Gly Leu Ala His Tyr
20 25 30
Arg Leu Gly Trp His Ala Leu Glu Gly Ile Phe Trp Trp Gly Ala Ser
35 40 45
Val Phe His Ala Leu Gln Ala Met Leu Val Arg Lys Trp Pro Leu Gly
50 55 60
Leu Val Glu Phe Thr Gly Thr Cys Gly Ile Leu Val Glu Val Ile Gly
65 70 75 80
Leu Trp Trp Gly Glu Gly Ser Thr Gly Asn Arg Trp Met Gly Leu Asn
85 90 95
Ser Thr Gly Gly Gln *
100 101

<210> 1784
<211> 243
<212> PRT
<213> Homo sapiens

<400> 1784
Met Gly Glu Ala Ser Pro Pro Ala Pro Ala Arg Arg His Leu Leu Val
1 5 10 15
Leu Leu Leu Leu Leu Ser Thr Leu Val Ile Pro Ser Ala Ala Ala Pro
20 25 30
Ile His Asp Ala Asp Ala Gln Glu Ser Ser Leu Gly Leu Thr Gly Leu
35 40 45
Gln Ser Leu Leu Gln Gly Phe Ser Arg Leu Phe Leu Lys Gly Asn Leu
50 55 60
Leu Arg Gly Ile Asp Ser Leu Phe Ser Ala Pro Met Asp Phe Arg Gly
65 70 75 80
Leu Pro Gly Asn Tyr His Lys Glu Glu Asn Gln Glu His Gln Leu Gly
85 90 95
Asn Asn Thr Leu Ser Ser His Leu Gln Ile Asp Lys Met Thr Asp Asn
100 105 110
Lys Thr Gly Glu Val Leu Ile Ser Glu Asn Val Val Ala Ser Ile Gln
115 120 125
Pro Ala Glu Gly Ser Phe Glu Gly Asp Leu Lys Val Pro Arg Met Glu
130 135 140
Glu Lys Glu Ala Leu Val Pro Ile Gln Lys Ala Thr Asp Ser Phe His
145 150 155 160
Thr Glu Leu His Pro Arg Val Ala Phe Trp Ile Ile Lys Leu Pro Arg
165 170 175
Arg Arg Ser His Gln Asp Ala Leu Glu Gly Gly His Trp Leu Ser Glu
180 185 190
Lys Arg His Arg Leu Gln Ala Ile Arg Asp Gly Leu Arg Lys Gly Thr
195 200 205
His Lys Asp Val Leu Glu Glu Gly Thr Glu Ser Ser Ser His Ser Arg
210 215 220
Leu Ser Pro Arg Lys Thr His Leu Leu Tyr Ile Leu Arg Pro Ser Arg
225 230 235 240
Gln Leu *
242

```
<210> 1785
<211> 158
<212> PRT
<213> Homo sapiens

<400> 1785
Met Lys Ala Leu Leu Leu Leu Val Leu Pro Trp Leu Ser Pro Ala Asn
1               5                   10                  15
Tyr Ile Asp Asn Val Gly Asn Leu His Phe Leu Tyr Ser Glu Leu Cys
            20                  25                  30
Lys Gly Ala Ser His Tyr Gly Leu Thr Lys Asp Arg Lys Arg Arg Ser
        35                  40                  45
Gln Asp Gly Cys Pro Asp Gly Cys Ala Ser Leu Thr Ala Thr Ala Pro
    50                  55                  60
Ser Pro Glu Val Ser Ala Ala Ala Thr Ile Ser Leu Met Thr Asp Glu
65                  70                  75                  80
Pro Gly Leu Asp Asn Pro Ala Tyr Val Ser Ser Ala Glu Asp Gly Gln
                85                  90                  95
Pro Ala Ile Ser Pro Val Asp Ser Gly Arg Ser Asn Arg Thr Arg Ala
            100                 105                 110
Arg Pro Phe Glu Arg Ser Thr Ile Ile Ser Arg Ser Phe Lys Lys Ile
        115                 120                 125
Asn Arg Ala Leu Ser Val Leu Arg Arg Thr Lys Ser Gly Ser Ala Val
    130                 135                 140
Ala Asn His Ala Asp Gln Gly Arg Glu Asn Ser Glu Asn Thr
145                 150                 155         158


<210> 1786
<211> 142
<212> PRT
<213> Homo sapiens

<400> 1786
Met Glu Ser Ala Val Arg Val Glu Ser Gly Val Leu Val Gly Val Val
1               5                   10                  15
Cys Leu Leu Leu Ala Cys Pro Ala Thr Ala Thr Gly Pro Glu Val Ala
            20                  25                  30
Gln Pro Glu Val Asp Thr Thr Leu Gly Arg Val Arg Gly Arg Gln Val
        35                  40                  45
Gly Val Lys Gly Thr Asp Arg Leu Val Asn Val Phe Leu Gly Ile Pro
    50                  55                  60
Phe Ala Gln Pro Pro Leu Gly Pro Asp Arg Phe Ser Ala Pro His Pro
65                  70                  75                  80
Ala Gln Pro Trp Glu Gly Val Arg Asp Ala Ser Thr Ala Pro Pro Met
                85                  90                  95
Cys Leu Gln Asp Val Glu Ser Met Asn Ser Ser Arg Phe Val Leu Asn
            100                 105                 110
Gly Lys Gln Gln Ile Phe Ser Val Ser Glu Asp Cys Leu Val Leu Asn
        115                 120                 125
Val Tyr Ser Pro Ala Glu Val Pro Ala Gly Ser Gly Arg Pro
    130                 135                 140         142
```

<210> 1787
<211> 120
<212> PRT
<213> Homo sapiens

<221> misc_feature
<222> (1)...(120)
<223> Xaa = any amino acid or nothing

<400> 1787
Met Ala Leu Thr Gly Tyr Ser Trp Leu Leu Leu Ser Ala Thr Phe Leu
1 5 10 15
Asn Val Gly Ala Glu Ile Ser Ile Thr Leu Glu Pro Ala Gln Pro Ser
20 25 30
Glu Gly Asp Asn Val Thr Leu Val Val His Gly Leu Ser Gly Glu Leu
35 40 45
Leu Ala Tyr Ser Trp Tyr Ala Gly Pro Thr Leu Ser Val Ser Tyr Leu
50 55 60
Val Ala Ser Tyr Ile Val Ser Thr Gly Asp Glu Thr Pro Gly Pro Ala
65 70 75 80
His Thr Xaa Arg Glu Ala Val Arg Pro Asp Gly Ser Leu Asp Ile Gln
85 90 95
Gly Ile Leu Pro Arg His Ser Ser Thr Tyr Ile Leu Gln Thr Phe Asn
100 105 110
Arg Gln Leu Gln Thr Glu Val Gly
115 120

<210> 1788
<211> 68
<212> PRT
<213> Homo sapiens

<400> 1788
Met Ser Trp Leu Ala Asn Gly Val Cys Leu Tyr Glu Tyr Leu Phe Phe
1 5 10 15
Arg Cys Gly Phe Leu Ile Leu Gln Pro Cys Ser Phe Asp Ala Ser Leu
20 25 30
Thr Asp Glu Glu Ser Arg Lys Asn Trp Glu Glu Phe Gly Asn Pro Asp
35 40 45
Gly Pro Gln Gly Val Val Asn Asp Asp Phe Lys Ile Leu Ala Ile Trp
50 55 60
Tyr Ile Leu *
65 67

<210> 1789
<211> 133
<212> PRT
<213> Homo sapiens

<400> 1789
Met Ala Val Val Ile Arg Leu Leu Gly Leu Pro Phe Ile Ala Gly Pro
1 5 10 15

Val Asp Ile Arg His Phe Phe Thr Gly Leu Thr Ile Pro Asp Gly Gly
20 25 30
Val His Ile Ile Gly Gly Glu Ile Gly Glu Ala Phe Ile Ile Phe Ala
35 40 45
Thr Asp Glu Asp Ala Arg Arg Ala Ile Ser Arg Ser Gly Gly Phe Ile
50 55 60
Lys Asp Ser Ser Val Glu Leu Phe Leu Ser Ser Lys Ala Glu Met Gln
65 70 75 80
Lys Thr Ile Glu Met Lys Arg Thr Asp Arg Val Gly Arg Gly Arg Pro
85 90 95
Gly Ser Gly Thr Ser Gly Val Asp Ser Leu Ser Asn Phe Ile Glu Ser
100 105 110
Val Lys Glu Glu Ala Ser Asn Ser Gly Tyr Gly Ser Ser Ile Asn Gln
115 120 125
Asp Ala Gly Phe His
130 133

<210> 1790
<211> 82
<212> PRT
<213> Homo sapiens

<400> 1790
Met Ala Ala Trp Gly Phe Cys Phe Ala Val Ser Ala Leu Val Val Ala
1 5 10 15
Cys Glu Phe Thr Arg Leu His Gly Cys Leu Arg Leu Ser Trp Gly Asn
20 25 30
Phe Thr Ala Ala Phe Ala Met Leu Ala Thr Leu Leu Cys Ala Thr Ala
35 40 45
Ala Val Leu Tyr Pro Leu Tyr Phe Ala Arg Arg Glu Cys Pro Pro Glu
50 55 60
Pro Ala Gly Cys Ala Ala Arg Asp Phe Arg Leu Ala Ala Ser Val Phe
65 70 75 80
Ala Gly
82

<210> 1791
<211> 50
<212> PRT
<213> Homo sapiens

<400> 1791
Met His Ala Ser Glu Gly Leu Pro Ala Leu Pro Leu Leu Ala Leu Val
1 5 10 15
Ser His Ser His Ser Cys Pro Pro Leu Pro Leu Gln Pro His His Leu
20 25 30
Pro Ala Ile Leu Phe Phe Leu Val Gly His Gln Leu Met Lys Cys Ile
35 40 45
Arg *
49

<210> 1792
<211> 166
<212> PRT
<213> Homo sapiens

<221> misc_feature
<222> (1)...(166)
<223> Xaa = any amino acid or nothing

<400> 1792
Met Leu Leu Trp Leu Leu Leu Leu Ile Leu Thr Pro Gly Arg Glu Gln
1 5 10 15
Ser Gly Val Ala Pro Lys Ala Val Leu Leu Leu Asp Pro Pro Trp Ser
20 25 30
Thr Ala Phe Lys Gly Glu Lys Val Ala Leu Ile Cys Ser Ser Ile Ser
35 40 45
His Ser Leu Ala Gln Gly Asp Thr Tyr Trp Tyr His Asp Glu Lys Leu
50 55 60
Leu Lys Ile Lys His Asp Lys Ile Gln Ile Thr Glu Pro Gly Asn Tyr
65 70 75 80
Gln Cys Lys Thr Arg Gly Ser Ser Leu Ser Asp Ala Val His Val Glu
85 90 95
Phe Ser Pro Asp Trp Leu Ile Leu Gln Ala Leu His Pro Val Phe Glu
100 105 110
Gly Asp Asn Val Ile Leu Arg Cys Gln Gly Lys Asp Asn Lys Asn Thr
115 120 125
His His Lys Val Tyr Tyr Lys Asp Gly Lys Gln Xaa Ser Asn Ser Tyr
130 135 140
Asn Leu Glu Lys Asn Thr Val Asp Ser Val Ser Arg Asp Asn Ser Pro
145 150 155 160
Tyr Tyr Cys Ala Gly *
165

<210> 1793
<211> 146
<212> PRT
<213> Homo sapiens

<400> 1793
Met Ala Thr Ala Ala Gln Gly Pro Leu Ser Leu Leu Trp Gly Trp Leu
1 5 10 15
Trp Ser Glu Arg Phe Trp Leu Pro Glu Asn Val Ser Trp Ala Asp Leu
20 25 30
Glu Gly Pro Ala Asp Gly Tyr Gly Tyr Pro Arg Gly Arg His Ile Leu
35 40 45
Ser Val Phe Pro Leu Ala Ala Gly Ile Phe Phe Val Arg Leu Leu Phe
50 55 60
Glu Arg Phe Ile Ala Lys Pro Cys Ala Leu Arg Ile Gly Ile Glu Asp
65 70 75 80
Ser Gly Pro Tyr Gln Ala Gln Pro Asn Ala Ile Leu Glu Lys Val Phe
85 90 95
Ile Ser Ile Thr Lys Tyr Pro Asp Lys Lys Arg Leu Glu Gly Leu Ser
100 105 110
Lys Gln Leu Asp Trp Asn Val Arg Lys Ile Gln Cys Trp Phe Arg His
115 120 125

Arg Arg Asn Gln Asp Lys Pro Pro Thr Leu Thr Lys Phe Cys Glu Ser
130 135 140
Met *
145

<210> 1794
<211> 151
<212> PRT
<213> Homo sapiens

<400> 1794
Met Glu Arg Arg Arg Leu Leu Gly Gly Met Ala Leu Leu Leu Leu Gln
1 5 10 15
Ala Leu Pro Ser Pro Leu Ser Ala Arg Ala Glu Pro Pro Gln Asp Lys
20 25 30
Glu Ala Cys Val Gly Thr Asn Asn Gln Ser Tyr Ile Cys Asp Thr Gly
35 40 45
His Cys Cys Gly Gln Ser Gln Cys Cys Asn Tyr Tyr Tyr Glu Leu Trp
50 55 60
Trp Phe Trp Leu Val Trp Thr Ile Ile Ile Ile Leu Ser Cys Cys Cys
65 70 75 80
Val Cys His His Arg Arg Ala Lys His Arg Leu Gln Ala Gln Gln Arg
85 90 95
Gln His Glu Ile Asn Leu Ile Ala Tyr Arg Glu Ala His Asn Tyr Ser
100 105 110
Ala Leu Pro Phe Tyr Phe Arg Phe Leu Pro Asn Tyr Leu Leu Pro Pro
115 120 125
Tyr Glu Glu Val Val Asn Arg Pro Pro Thr Pro Pro Pro Pro Tyr Ser
130 135 140
Ala Phe Gln Leu Gln Gln Gln
145 150 151

<210> 1795
<211> 177
<212> PRT
<213> Homo sapiens

<400> 1795
Met Ala Ala Leu Ala Ala Ala Ala Lys Lys Val Trp Ser Ala Arg Arg
1 5 10 15
Leu Leu Val Leu Leu Phe Thr Pro Leu Ala Leu Leu Pro Val Val Phe
20 25 30
Ala Leu Pro Pro Lys Glu Gly Arg Cys Leu Phe Val Ile Leu Leu Met
35 40 45
Ala Val Tyr Trp Cys Thr Glu Ala Leu Pro Leu Ser Val Thr Ala Leu
50 55 60
Leu Pro Ile Val Leu Phe Pro Phe Met Gly Ile Leu Pro Ser Asn Lys
65 70 75 80
Val Cys Pro Gln Tyr Phe Leu Asp Thr Asn Phe Leu Phe Leu Ser Gly
85 90 95
Leu Ile Met Ala Ser Ala Ile Glu Glu Trp Asn Leu His Arg Arg Ile
100 105 110
Ala Leu Lys Ile Leu Met Leu Val Gly Val Gln Pro Ala Arg Leu Ile

115 120 125
Leu Gly Met Met Val Thr Thr Ser Phe Leu Ser Met Trp Leu Ser Asn
130 135 140
Thr Ala Ser Thr Ala Met Met Leu Pro Ile Ala Asn Ala Ile Leu Lys
145 150 155 160
Ser Leu Phe Gly Gln Lys Glu Val Arg Lys Asp Pro Gln Pro Gly Glu
165 170 175 176
*

<210> 1796
<211> 98
<212> PRT
<213> Homo sapiens

<221> misc_feature
<222> (1)...(98)
<223> Xaa = any amino acid or nothing

<400> 1796
Met His Pro Leu Pro Gly Tyr Trp Ser Cys Tyr Cys Leu Leu Leu Leu
1 5 10 15
Phe Ser Leu Gly Val Gln Gly Ser Leu Gly Ala Pro Ser Ala Ala Pro
20 25 30
Glu Gln Val His Leu Ser Tyr Pro Gly Glu Pro Gly Ser Met Thr Val
35 40 45
Thr Trp Thr Thr Trp Val Pro Thr Arg Ser Glu Val Gln Phe Gly Leu
50 55 60
Gln Pro Ser Gly Pro Leu Pro Leu Arg Ala Gln Gly Thr Phe Val Pro
65 70 75 80
Phe Val Asp Xaa Gly Ile Leu Arg Arg Lys Leu Tyr Ile His Arg Val
85 90 95
Thr Leu
98

<210> 1797
<211> 96
<212> PRT
<213> Homo sapiens

<400> 1797
Met Phe Leu Trp Leu Phe Leu Ile Leu Ser Ala Leu Ile Ser Ser Thr
1 5 10 15
Asn Ala Asp Ser Asp Ile Ser Val Glu Ile Cys Asn Val Cys Ser Cys
20 25 30
Val Ser Val Glu Asn Val Leu Tyr Val Asn Cys Glu Lys Val Ser Val
35 40 45
Tyr Arg Pro Asn Gln Leu Lys Pro Pro Trp Ser Asn Phe Tyr His Leu
50 55 60
Asn Phe Gln Asn Asn Phe Leu Asn Ile Leu Tyr Pro Asn Thr Phe Leu
65 70 75 80
Asn Phe Ser His Ala Val Ser Leu His Leu Gly Asn Asn Lys Leu Gln
85 90 95 96

<210> 1798
<211> 91
<212> PRT
<213> Homo sapiens

<400> 1798
```
Met Arg Pro Ala Leu Ala Val Gly Leu Val Phe Ala Gly Cys Cys Ser
 1               5                  10                  15
Asn Val Ile Phe Leu Glu Leu Leu Ala Arg Lys His Pro Gly Cys Gly
            20                  25                  30
Asn Ile Val Thr Phe Ala Gln Phe Leu Phe Ile Ala Val Glu Gly Phe
        35                  40                  45
Leu Phe Glu Ala Asp Leu Gly Arg Lys Pro Pro Ala Ile Pro Ile Arg
    50                  55                  60
Tyr Tyr Ala Ile Met Val Thr Met Phe Phe Thr Val Ser Val Val Asn
65                  70                  75                  80
Asn Tyr Ala Leu Asn Leu Asn Ile Ala Met Pro
                85                  90  91
```

<210> 1799
<211> 77
<212> PRT
<213> Homo sapiens

<400> 1799
```
Met Arg Ser Leu Val Trp Val Leu Ile Gln Gln Leu Thr Pro Leu Tyr
 1               5                  10                  15
Lys Gly Glu Thr Trp Thr Gln Thr Cys Thr Glu Asp His Val Thr Met
            20                  25                  30
Lys Ala Glu Ile Arg Val Met Leu Leu Glu Ala Arg Glu Asp Cys Gln
        35                  40                  45
Leu Met Thr Lys Arg Ser Gln Glu Thr Gly Leu Gln Arg Ile Leu Pro
    50                  55                  60
Glu Gly Ser Gln Lys Glu Pro Thr Leu Thr Thr Pro  *
65                  70                  75  76
```

<210> 1800
<211> 182
<212> PRT
<213> Homo sapiens

<400> 1800
```
Met Ser Leu Lys Met Leu Ile Ser Arg Asn Lys Leu Ile Leu Leu Leu
 1               5                  10                  15
Gly Ile Val Phe Phe Glu Arg Gly Lys Ser Ala Thr Leu Ser Leu Pro
            20                  25                  30
Lys Ala Pro Ser Cys Gly Gln Ser Leu Val Lys Val Gln Pro Trp Asn
```

```
        35                  40                  45
Tyr Phe Asn Ile Phe Ser Arg Ile Leu Gly Gly Ser Gln Val Glu Lys
    50                  55                  60
Gly Ser Tyr Pro Trp Gln Val Ser Leu Lys Gln Arg Gln Lys His Ile
65                  70                  75                  80
Cys Gly Gly Ser Ile Val Ser Pro Gln Trp Val Ile Thr Ala Ala His
                85                  90                  95
Cys Ile Ala Asn Arg Asn Ile Val Ser Thr Leu Asn Val Thr Ala Gly
            100                 105                 110
Glu Tyr Asp Leu Ser Gln Thr Asp Pro Gly Glu Gln Thr Leu Thr Ile
        115                 120                 125
Glu Thr Val Ile Ile His Pro His Phe Ser Thr Lys Lys Pro Met Asp
    130                 135                 140
Tyr Asp Ile Ala Leu Leu Lys Met Ala Gly Ala Phe Gln Phe Gly His
145                 150                 155                 160
Phe Val Gly Pro Ile Cys Leu Pro Glu Leu Arg Glu Gln Phe Glu Ala
                165                 170                 175
Gly Phe Ile Cys Thr Thr
            180     182
```

```
<210> 1801
<211> 202
<212> PRT
<213> Homo sapiens
```

```
<400> 1801
Met Thr Glu Ala Thr Phe Asp Thr Leu Arg Leu Trp Leu Ile Ile Leu
1               5                   10                  15
Leu Cys Ala Leu Arg Leu Ala Met Met Arg Ser His Leu Gln Ala Tyr
            20                  25                  30
Leu Asn Leu Ala Gln Lys Cys Val Asp Gln Met Lys Lys Glu Ala Gly
        35                  40                  45
Arg Ile Ser Thr Val Glu Leu Gln Lys Met Val Ala Arg Val Phe Tyr
    50                  55                  60
Tyr Leu Cys Val Ile Ala Leu Gln Tyr Val Ala Pro Leu Val Met Leu
65                  70                  75                  80
Leu His Thr Thr Leu Leu Leu Lys Thr Leu Gly Asn His Ser Trp Gly
                85                  90                  95
Ile Tyr Pro Glu Ser Ile Ser Thr Leu Pro Val Asp Asn Ser Leu Leu
            100                 105                 110
Ser Asn Ser Val Tyr Ser Glu Leu Pro Ser Ala Glu Gly Lys Met Lys
        115                 120                 125
His Asn Ala Arg Gln Gly Pro Ala Val Pro Pro Gly Met Gln Ala Tyr
    130                 135                 140
Gly Ala Ala Pro Phe Glu Asp Leu Gln Leu Asp Phe Thr Glu Met Pro
145                 150                 155                 160
Lys Cys Gly Asp Leu Ile Pro Arg Phe Gly Leu Pro Leu Arg Ile Gly
                165                 170                 175
Ser Asp Asn Gly Leu Ala Phe Val Ala Asp Leu Val Gln Lys Thr Ala
            180                 185                 190
Lys Trp Lys Gly Pro Gln Ile Val Val Leu
        195                 200     202
```

```
<210> 1802
```

<211> 172
<212> PRT
<213> Homo sapiens

<400> 1802

```
Met Asn Asn Phe Arg Ala Thr Ile Leu Phe Trp Ala Ala Ala Ala Trp
 1               5                  10                  15
Ala Lys Ser Gly Lys Pro Ser Gly Glu Met Asp Glu Val Gly Val Gln
            20                  25                  30
Lys Cys Lys Asn Ala Leu Lys Leu Pro Val Leu Glu Val Leu Pro Gly
        35                  40                  45
Gly Gly Trp Asp Asn Leu Arg Asn Val Asp Met Gly Arg Val Met Glu
    50                  55                  60
Leu Thr Tyr Ser Asn Cys Arg Thr Thr Glu Asp Gly Gln Tyr Ile Ile
65                  70                  75                  80
Pro Asp Glu Ile Phe Thr Ile Pro Gln Lys Gln Ser Asn Leu Glu Met
                85                  90                  95
Asn Ser Glu Ile Leu Glu Ser Trp Ala Asn Tyr Gln Ser Ser Thr Ser
            100                 105                 110
Tyr Ser Ile Asn Thr Glu Leu Ser Leu Phe Ser Lys Val Asn Gly Lys
        115                 120                 125
Phe Ser Thr Glu Phe Gln Arg Met Lys Thr Leu Gln Val Lys Asp Gln
    130                 135                 140
Ala Ile Thr Thr Arg Val Gln Val Arg Asn Leu Val Tyr Thr Val Lys
145                 150                 155                 160
Ile Asn Pro Thr Leu Glu Leu Ser Ser Gly Phe Arg
                165                 170         172
```

<210> 1803
<211> 158
<212> PRT
<213> Homo sapiens

<400> 1803

```
Met Ser Leu Arg Leu Gly Pro Ala Trp Arg His Leu Thr Cys Leu Gly
 1               5                  10                  15
Thr Lys His Ser Lys Ala Asn Ser Val Leu Ala Ser Gln His Ala Gly
            20                  25                  30
Phe Phe Val Ala Gln Gly Arg Trp Ala Ile His Arg Ala Phe Ser Ser
        35                  40                  45
Arg Thr Ser Pro Thr Pro Pro Arg Gly Pro Leu Leu Leu Pro Gly Arg
    50                  55                  60
His Pro Leu Leu Ser Arg Arg Arg Ala Gln Ala Ile Arg Ser Ser Thr
65                  70                  75                  80
Arg Pro Ser Leu Pro Ala His Leu Phe Lys Pro Ala Pro Ala Ile Ala
                85                  90                  95
Leu Ile Val Ser Pro Leu Arg Phe Pro Arg Arg Thr Ser Pro Cys His
            100                 105                 110
Leu Ser Gly Pro Pro Ala Pro Pro Cys Arg Thr Leu His Thr Leu Leu
        115                 120                 125
Arg Pro Val Cys Val Val Arg Arg Thr Pro Pro Val Phe Phe Thr Ser
    130                 135                 140
Phe Thr Pro Ala Arg Ala Ala Val Ala Ser His Pro Thr Pro
145                 150                 155             158
```

<210> 1804
<211> 102
<212> PRT
<213> Homo sapiens

<400> 1804
Met Gly Leu Gly Gln Pro Gln Ala Trp Leu Leu Gly Leu Pro Thr Ala
1 5 10 15
Val Val Tyr Gly Ser Leu Ala Leu Phe Thr Thr Ile Leu His Asn Val
20 25 30
Phe Leu Leu Tyr Tyr Val Asp Thr Phe Val Ser Val Tyr Lys Ile Asn
35 40 45
Lys Met Ala Phe Trp Val Gly Glu Thr Val Phe Leu Leu Trp Asn Ser
50 55 60
Leu Asn Asp Pro Leu Phe Gly Trp Leu Ser Asp Arg Gln Phe Leu Ser
65 70 75 80
Ser Gln Pro Arg Ser Gly Ala Gly Leu Ser Ser Arg Ala Val Val Leu
85 90 95
Ala Arg Val Gln Ala Leu
100 102

<210> 1805
<211> 54
<212> PRT
<213> Homo sapiens

<400> 1805
Met Ala Asp Ser Val Leu Thr Leu Val Phe Thr Ser Cys Leu Leu Ser
1 5 10 15
Glu Leu Ser Leu Val Cys Ser Asp Phe Arg Pro Thr Pro Ile Ser Tyr
20 25 30
Gln Ser Arg Tyr Gly Ser Gly Asp Gly Trp Ile Arg Cys Lys Ser Glu
35 40 45
Val Arg Glu Thr Gln *
50 53

<210> 1806
<211> 56
<212> PRT
<213> Homo sapiens

<400> 1806
Met Leu Ser Val Lys Arg Phe Arg Ala Met Val Met Phe Phe Met Ala
1 5 10 15
Met Val Ala Met Met Lys Asn Lys Cys Gln Gln Thr Asn Glu Ala Lys
20 25 30
Phe Cys Val His Met Tyr Leu His Phe Tyr Phe Ser Ser His Ser Ser
35 40 45
Ala Val Cys Ile Ser Ser Pro Leu
50 55 56

<210> 1807
<211> 47
<212> PRT
<213> Homo sapiens

<400> 1807
Met Gln Ser Met Ile Asn Met Ile Val Ser Leu Leu Gly Leu Val Ala
1 5 10 15
Thr Val Thr Leu Ile Pro Ala Phe Arg Gly His Phe Ile Ala Ala Arg
20 25 30
Leu Gly Gly Gln Ser Leu Gly Lys Thr Ser Arg Gln His Met *
35 40 45 46

<210> 1808
<211> 119
<212> PRT
<213> Homo sapiens

<400> 1808
Met Ala Ala Ser Leu Leu Ala Val Leu Leu Leu Leu Leu Leu Glu Arg
1 5 10 15
Gly Met Phe Ser Ser Pro Ser Pro Pro Pro Ala Leu Leu Glu Lys Val
20 25 30
Phe Gln Tyr Ile Asp Leu His Gln Asp Glu Phe Val Gln Thr Leu Lys
35 40 45
Glu Trp Val Ala Ile Glu Ser Asp Ser Val Gln Pro Val Pro Arg Phe
50 55 60
Arg Gln Glu Leu Phe Arg Met Met Ala Val Ala Ala Asp Thr Leu Gln
65 70 75 80
Arg Leu Gly Ala Arg Val Ala Ser Val Asp Met Gly Pro Gln Gln Leu
85 90 95
Pro Asp Gly Gln Ser Leu Pro Ile Pro Pro Val Ile Leu Ala Glu Leu
100 105 110
Gly Ser Asp Pro Thr Lys Gly
115 119

<210> 1809
<211> 91
<212> PRT
<213> Homo sapiens

<400> 1809
Met Ser Arg Ser His Val Ala Leu Leu Gly Leu Ser Leu Leu Leu Met
1 5 10 15
Leu Leu Leu Tyr Ala Gly Leu Pro Ser Pro Pro Glu Gln Thr Ser Cys
20 25 30
Leu Trp Gly Asp Pro Asn Val Thr Val Leu Ala Val Ser Thr Pro Ala
35 40 45
Asn Ser Pro Met Phe Tyr Leu Glu Gly Leu Pro Leu His Leu Ala His

```
    50                  55                  60
Arg Val Asp Val Ile Pro Leu Ser Ser Leu Gly Pro Leu Val Ser Pro
 65                 70                  75                  80
Leu Arg Cys Gln Ala Leu Pro Pro Arg Leu Ser
                85                  90  91
```

<210> 1810
<211> 58
<212> PRT
<213> Homo sapiens

<400> 1810

```
Met Leu Leu Phe Gly Leu Cys Trp Gly Pro Tyr Val Ala Thr Leu Leu
  1               5                  10                  15
Leu Ser Val Leu Ala Tyr Glu Gln Arg Pro Pro Leu Gly Pro Gly Thr
            20                  25                  30
Leu Leu Ser Leu Leu Ser Leu Gly Ser Ala Lys Ala Ala Ala Val Pro
        35                  40                  45
Val Ala Met Gly Leu Gly Asp Gln Arg Tyr
    50                  55          58
```

<210> 1811
<211> 48
<212> PRT
<213> Homo sapiens

<400> 1811

```
Met Ala Ser Ala Ser Phe Ser Leu Leu Ile Cys Gly Phe Leu Ala Ser
  1               5                  10                  15
Leu Ser Leu Gln Arg Ile Glu Glu Leu Gly Leu Gly Leu Gly Leu Gly
            20                  25                  30
Phe Gly Leu Arg Glu Cys Cys Gly Trp Phe Gly Leu Leu Ser Leu Val
        35                  40                  45          48
```

<210> 1812
<211> 84
<212> PRT
<213> Homo sapiens

<400> 1812

```
Met Lys Val Leu Leu Ala Val Ala Leu Ile Ala Arg Thr Val Phe Phe
  1               5                  10                  15
Leu Leu Leu Ala Gly Pro Ser Ala Ala Asp Asp Lys Lys Lys Gly Pro
            20                  25                  30
Lys Val Thr Val Lys Val Tyr Phe Asp Leu Arg Ile Gly Asp Glu Asp
        35                  40                  45
Val Arg Arg Glu Ile Phe Gly Leu Phe Gly Lys Thr Ala Pro Lys Thr
    50                  55                  60
```

Glu Asp Asn Phe Val Ala Leu Ala Thr Gly Gln Lys Gly Phe Gly Tyr
65 70 75 80
Lys Asn Ser *
83

<210> 1813
<211> 46
<212> PRT
<213> Homo sapiens

<400> 1813
Met Ala Ala Ala Asp Asp Thr Ile Leu Gly Phe Arg Ala Ala Leu Leu
1 5 10 15
Ile Leu Val Ala Ala Ala Ala Ala Leu Ser Pro Lys Val Ala Cys Arg
20 25 30
Val Gly Thr Val Arg Arg Arg Glu Thr Pro Gln Pro Ser Ala
35 40 45 46

<210> 1814
<211> 65
<212> PRT
<213> Homo sapiens

<400> 1814
Met Ile Ile Tyr Leu Thr Phe Pro Val Ala Met Phe Trp Val Ser Asn
1 5 10 15
Gln Ala Glu Trp Phe Glu Asp Asp Val Ile Gln Arg Lys Arg Glu Leu
20 25 30
Trp Pro Pro Glu Lys Leu Gln Glu Ile Glu Glu Phe Lys Glu Arg Leu
35 40 45
Arg Lys Arg Arg Glu Glu Lys Leu Leu Arg Asp Ala Gln Gln Asn Ser
50 55 60 64
*

<210> 1815
<211> 100
<212> PRT
<213> Homo sapiens

<400> 1815
Met Phe Lys Ser Lys Leu Leu Asn Phe Tyr Ile Phe Val Asn Cys Met
1 5 10 15
Asn Phe Leu Met Leu Ser Ile Ala Ser Phe Asn Pro Phe Trp Ser Glu
20 25 30
Ile Ile Val Cys Asn Ile Gln Phe Phe Tyr Tyr Thr Leu Ser Ser Arg
35 40 45
Val His Val Gln Asn Val Gln Val Cys Tyr Thr Gly Ile His Val Pro
50 55 60
Cys Trp Phe Ala Ala Pro Ile Asn Ser Ser Phe Thr Leu Gly Ile Ser

```
 65                  70                  75                  80
Pro Asn Ala Ile Pro Phe Ile Val Pro His Pro Gln Thr Gly Pro Asn
                 85                  90                  95
Val Arg Cys Ser
            100
```

```
     <210> 1816
     <211> 115
     <212> PRT
     <213> Homo sapiens

     <221> misc_feature
     <222> (1)...(115)
     <223> Xaa = any amino acid or nothing
```

```
     <400> 1816
Met Phe Cys Phe Leu Val Ser Val Leu Tyr Ser Lys Ala Lys Leu Ala
  1               5                  10                  15
Ser Ala Cys Gly Gly Ile Ile Tyr Phe Leu Ser Tyr Val Pro Tyr Met
             20                  25                  30
Tyr Val Ala Ile Arg Glu Glu Val Ala His Asp Lys Ile Thr Ala Phe
         35                  40                  45
Glu Lys Cys Ile Ala Ser Leu Met Ser Thr Thr Ala Phe Gly Leu Gly
     50                  55                  60
Ser Lys Tyr Phe Ala Leu Tyr Glu Val Pro Gly Val Gly Ile Gln Trp
 65                  70                  75                  80
His Thr Phe Ser Gln Ser Pro Val Glu Gly Glu Asp Leu Asn Leu Pro
                 85                  90                  95
Pro Pro Pro Pro Met Met Pro Ala Pro Xaa Val Val Tyr Gly Ile Leu
            100                 105                 110
Thr Lys  *
    114
```

```
     <210> 1817
     <211> 144
     <212> PRT
     <213> Homo sapiens
```

```
     <400> 1817
Met Val Leu Gly Leu Leu Val Gln Ile Trp Ala Leu Gln Glu Ala Ser
  1               5                  10                  15
Ser Leu Ser Val Gln Gln Gly Pro Asn Leu Leu Gln Val Arg Gln Gly
             20                  25                  30
Ser Gln Ala Thr Leu Val Cys Gln Val Asp Gln Ala Thr Ala Trp Glu
         35                  40                  45
Arg Leu Arg Val Lys Trp Thr Lys Asp Gly Ala Ile Leu Cys Gln Pro
     50                  55                  60
Tyr Ile Thr Asn Gly Ser Leu Ser Leu Gly Val Cys Gly Pro Gln Gly
 65                  70                  75                  80
Arg Leu Ser Trp Gln Ala Pro Ser His Leu Thr Leu Gln Leu Asp Pro
                 85                  90                  95
Val Ser Leu Asn His Ser Gly Ala Tyr Val Cys Trp Ala Ala Val Glu
            100                 105                 110
```

Ile Pro Glu Leu Glu Glu Ala Glu Gly Asn Ile Thr Arg Leu Phe Val
115 120 125
Asp Pro Asp Asp Pro Thr Gln Asn Arg Asn Arg Ile Ala Ser Phe Pro
130 135 140 144

<210> 1818
<211> 115
<212> PRT
<213> Homo sapiens

<400> 1818
Met Gln Ala Asp Arg Gly Gly Val Leu Phe Leu Val Ala Leu Pro Gly
1 5 10 15
Leu Trp Glu Thr Val Leu Arg His Pro Gly Ala Ser Pro Glu Pro Val
20 25 30
Ser Leu His Thr Gly Leu Ala Ala Glu Pro Leu Leu Gly Trp Arg Ala
35 40 45
Glu Val Ala Thr Ala Ala Gly Leu Gln Asp Arg Arg Ile Gly Arg Arg
50 55 60
Ser Leu Pro Ala Thr Leu Pro Pro Pro Phe Pro Gln Ala Gly Asp Leu
65 70 75 80
Arg Glu Ser Ile Leu Leu Leu Pro Cys Arg Glu Ser Arg Ser Thr Ser
85 90 95
Trp Leu Ser Pro Tyr Trp Val Pro Glu Ile Pro Gly Thr Leu His Asp
100 105 110
Arg Gly Arg
115

<210> 1819
<211> 70
<212> PRT
<213> Homo sapiens

<400> 1819
Met Pro Trp Leu Leu Ser Ala Pro Lys Leu Val Pro Ala Val Ala Asn
1 5 10 15
Val Arg Gly Leu Ser Gly Cys Met Leu Cys Ser Gln Arg Arg Tyr Ser
20 25 30
Leu Gln Pro Val Pro Glu Arg Arg Ile Pro Asn Arg Tyr Leu Gly Gln
35 40 45
Pro Ser Pro Phe Thr His Pro His Leu Leu Arg Pro Asp Ser Asn Ser
50 55 60
Cys Trp Glu Val Gly *
65 69

<210> 1820
<211> 635
<212> PRT
<213> Homo sapiens

<400> 1820
Met Leu Arg Ser Leu Leu Val Tyr Met Leu Phe Leu Leu Val Thr Leu
1 5 10 15
Leu Ala Ser Tyr Gly Asp Ala Ser Cys His Gly His Ala Tyr Arg Leu
20 25 30
Gln Ser Ala Ile Lys Gln Glu Leu His Ser Arg Ala Phe Leu Ala Ile
35 40 45
Thr Arg Ser Glu Glu Leu Trp Pro Trp Met Ala His Val Leu Leu Pro
50 55 60
Tyr Val His Gly Asn Gln Ser Ser Pro Glu Leu Gly Pro Pro Arg Leu
65 70 75 80
Arg Gln Val Arg Leu Gln Glu Ala Leu Tyr Pro Asp Pro Pro Gly Pro
85 90 95
Arg Val His Thr Cys Ser Ala Ala Gly Gly Phe Ser Thr Ser Asp Tyr
100 105 110
Asp Val Gly Trp Glu Ser Pro His Asn Gly Ser Gly Thr Trp Ala Tyr
115 120 125
Ser Ala Pro Asp Leu Leu Gly Ala Trp Ser Trp Gly Ser Cys Ala Val
130 135 140
Tyr Asp Ser Gly Gly Tyr Val Gln Glu Leu Gly Leu Ser Leu Glu Glu
145 150 155 160
Ser Arg Asp Arg Leu Arg Phe Leu Gln Leu His Asn Trp Leu Asp Asn
165 170 175
Arg Ser Arg Ala Val Phe Leu Glu Leu Thr Arg Tyr Ser Pro Ala Val
180 185 190
Gly Leu His Ala Ala Val Thr Leu Arg Leu Glu Phe Pro Ala Ala Gly
195 200 205
Arg Ala Leu Ala Ala Leu Ser Val Arg Pro Phe Ala Leu Arg Arg Leu
210 215 220
Ser Ala Gly Leu Ser Leu Pro Leu Leu Thr Ser Val Cys Leu Leu Leu
225 230 235 240
Phe Ala Val His Phe Ala Val Ala Glu Ala Arg Thr Trp His Arg Glu
245 250 255
Gly Arg Trp Arg Val Leu Arg Leu Gly Ala Trp Ala Arg Trp Leu Leu
260 265 270
Val Ala Leu Thr Ala Ala Thr Ala Leu Val Arg Leu Ala Gln Leu Gly
275 280 285
Ala Ala Asp Arg Gln Trp Thr Arg Phe Val Arg Gly Arg Pro Arg Arg
290 295 300
Phe Thr Ser Phe Asp Gln Val Ala His Val Ser Ser Ala Ala Arg Gly
305 310 315 320
Leu Ala Ala Ser Leu Leu Phe Leu Leu Leu Val Lys Ala Ala Gln His
325 330 335
Val Arg Phe Val Arg Gln Trp Ser Val Phe Gly Lys Thr Leu Cys Arg
340 345 350
Ala Leu Pro Glu Leu Leu Gly Val Thr Leu Gly Leu Val Val Leu Gly
355 360 365
Val Ala Tyr Ala Gln Leu Ala Ile Leu Leu Val Ser Ser Cys Val Asp
370 375 380
Ser Leu Trp Ser Val Ala Gln Ala Leu Leu Val Leu Cys Pro Gly Thr
385 390 395 400
Gly Leu Ser Thr Leu Cys Pro Ala Glu Ser Trp His Leu Ser Pro Leu
405 410 415
Leu Cys Val Gly Leu Trp Ala Leu Arg Leu Trp Gly Ala Leu Arg Leu
420 425 430
Gly Ala Val Ile Leu Arg Trp Arg Tyr His Ala Leu Arg Gly Glu Leu
435 440 445

```
Tyr Arg Pro Ala Trp Glu Pro Gln Asp Tyr Glu Met Val Glu Leu Phe
    450                 455                 460
Leu Arg Arg Leu Arg Leu Trp Met Gly Leu Ser Lys Val Lys Glu Phe
465                 470                 475                 480
Arg His Lys Val Arg Phe Glu Gly Met Glu Pro Leu Pro Ser Arg Ser
                485                 490                 495
Ser Arg Gly Ser Lys Val Ser Pro Asp Val Pro Pro Pro Ser Ala Gly
            500                 505                 510
Ser Asp Ala Ser His Pro Ser Thr Ser Ser Ser Gln Leu Asp Gly Leu
        515                 520                 525
Ser Val Ser Leu Gly Arg Leu Gly Thr Arg Cys Glu Pro Glu Pro Ser
    530                 535                 540
Arg Leu Gln Ala Val Phe Glu Ala Leu Leu Thr Gln Phe Asp Arg Leu
545                 550                 555                 560
Asn Gln Ala Thr Glu Asp Val Tyr Gln Leu Glu Gln Gln Leu His Ser
                565                 570                 575
Leu Gln Gly Arg Arg Ser Ser Arg Ala Pro Ala Gly Ser Ser Arg Gly
            580                 585                 590
Pro Ser Pro Gly Leu Arg Pro Ala Leu Pro Ser Arg Leu Ala Arg Ala
        595                 600                 605
Ser Arg Gly Val Asp Leu Ala Thr Gly Pro Ser Arg Thr Pro Leu Arg
    610                 615                 620
Ala Lys Asn Lys Val His Pro Ser Ser Thr  *
625                 630             634
```

<210> 1821
<211> 84
<212> PRT
<213> Homo sapiens

<400> 1821

```
Met Gly Ser Thr Trp Gly Ser Pro Gly Trp Val Arg Leu Ala Leu Cys
 1               5                  10                  15
Leu Thr Gly Leu Met Leu Ser Leu Tyr Thr Leu His Val Lys Ala Ala
            20                  25                  30
Arg Ala Arg Asn Arg Asp Tyr Arg Ala Leu Cys Asp Val Gly Thr Val
        35                  40                  45
Ile Ser Cys Thr Arg Val Phe Tyr Ser Lys Leu Pro Ala Asp Thr Leu
    50                  55                  60
Asp Leu Cys Pro Asp Ala Ala Glu Leu Pro Gly Val Ser Arg Trp Phe
65                  70                  75                  80
Cys Leu Pro Gly
            84
```

<210> 1822
<211> 108
<212> PRT
<213> Homo sapiens

<400> 1822

```
Met Ala Leu Asp Phe Val Asn Val Leu Leu Cys Gln Leu Ala Glu Val
 1               5                  10                  15
Thr Leu Gly Val Leu Arg Glu Glu Gly Ala Ser Leu Leu Val Ala Leu
```

```
            20                       25                       30
Gly Ser Ala Leu Phe Pro Ser Ala Ala Ala Val Gly Lys Gln Gly Ser
        35                       40                       45
Met Gly Val Thr Ser His Met Gln Cys Pro Val Cys Gln His Pro Arg
    50                       55                       60
Asp Val Leu Leu Ala Ser Pro Val Ser His Ser His Ala Cys Gln Pro
65                       70                       75                  80
Gln Pro Ala Gly Cys Ser Asn Cys His Leu Gly His Leu Thr Arg Ser
                    85                       90                       95
Pro Pro Phe Gln Gly Leu Leu Pro Leu Leu Gln  *
            100                      105     107
```

<210> 1823
<211> 74
<212> PRT
<213> Homo sapiens

<400> 1823

```
Met Gly Val Val Leu Tyr Val Met Leu Cys Ala Ser Leu Pro Phe Asp
 1                   5                       10                      15
Asp Thr Asp Ile Pro Lys Met Leu Trp Gln Gln Gln Lys Gly Val Ser
            20                       25                       30
Phe Pro Thr His Leu Ser Ile Ser Ala Asp Cys Gln Asp Leu Leu Lys
        35                       40                       45
Arg Leu Leu Glu Pro Asp Met Ile Leu Arg Pro Ser Ile Glu Glu Val
    50                       55                       60
Ser Trp His Pro Trp Leu Ala Ser Thr  *
65                       70          73
```

<210> 1824
<211> 58
<212> PRT
<213> Homo sapiens

<400> 1824

```
Met Ser Leu Ser Cys Thr Gly Phe Ala Leu Glu Lys Arg Cys Ala Gly
 1                   5                       10                      15
Trp Val Trp Trp Leu Thr Pro Val Ile Pro Ala Leu Leu Gly Gly Gln
            20                       25                       30
Gly Arg Gln Ile Met Ile Met Val Arg Ser Leu Arg Pro Ala Gly Pro
        35                       40                       45
Thr Trp Gly Asn Leu Ser Thr Thr Lys Thr
    50                       55          58
```

<210> 1825
<211> 225
<212> PRT
<213> Homo sapiens

<400> 1825

```
Met Ala Cys Lys Gly Leu Leu Gln Gln Val Gln Gly Pro Arg Leu Pro
 1               5                  10                  15
Trp Thr Arg Leu Leu Leu Leu Leu Leu Val Phe Ala Val Gly Phe Leu
            20                  25                  30
Cys His Asp Leu Arg Ser His Ser Ser Phe Gln Ala Ser Leu Thr Gly
        35                  40                  45
Arg Leu Leu Arg Ser Ser Gly Phe Leu Pro Ala Ser Gln Gln Ala Cys
    50                  55                  60
Ala Lys Leu Tyr Ser Tyr Ser Leu Gln Gly Tyr Ser Trp Leu Gly Glu
65                  70                  75                  80
Thr Leu Pro Leu Trp Gly Ser His Leu Leu Thr Val Val Arg Pro Ser
                85                  90                  95
Leu Gln Leu Ala Trp Ala His Thr Asn Ala Thr Val Ser Phe Leu Ser
            100                 105                 110
Ala His Cys Ala Ser His Leu Ala Trp Phe Gly Asp Ser Leu Thr Ser
        115                 120                 125
Leu Ser Gln Arg Leu Gln Ile Gln Leu Pro Asp Ser Val Asn Gln Leu
    130                 135                 140
Leu Arg Tyr Leu Arg Glu Leu Pro Leu Leu Phe His Gln Asn Val Leu
145                 150                 155                 160
Leu Pro Leu Trp His Leu Leu Leu Glu Ala Leu Ala Trp Ala Gln Glu
                165                 170                 175
His Cys His Glu Ala Cys Arg Gly Glu Val Thr Trp Asp Cys Met Lys
            180                 185                 190
Thr Gln Leu Ser Glu Ala Val His Trp Thr Trp Leu Cys Leu Gln Asp
        195                 200                 205
Ile Thr Val Ala Phe Leu Asp Trp Ala Leu Ala Leu Ile Ser Gln Gln
    210                 215                 220                 224
*
```

```
<210> 1826
<211> 119
<212> PRT
<213> Homo sapiens
```

```
<400> 1826
Met Tyr Arg Glu Val Cys Ser Ile Arg Phe Leu Phe Thr Ala Val Ser
 1               5                  10                  15
Leu Leu Ser Leu Phe Leu Ser Ala Phe Trp Leu Gly Leu Leu Tyr Leu
            20                  25                  30
Val Ser Pro Leu Glu Asn Glu Pro Lys Glu Met Leu Thr Leu Ser Glu
        35                  40                  45
Tyr His Glu Arg Ala Arg Ser Gln Gly Gln Gln Leu Leu Gln Phe Gln
    50                  55                  60
Ala Glu Leu Asp Lys Leu His Lys Glu Ala Ser Leu Val Cys Gly Cys
65                  70                  75                  80
Pro Ser Leu Arg Glu Val Pro Ser Ser Ala Val Ser Arg Leu Glu Pro
                85                  90                  95
Pro Ser Ile Ala Gln Pro Leu Leu Ser Arg Leu Gln Leu Tyr Leu Ser
            100                 105                 110
Asp Pro Ser Ser Tyr Leu Val
        115                 119
```

<210> 1827
<211> 58
<212> PRT
<213> Homo sapiens

<400> 1827
Met Lys Leu Met Arg Pro Leu Met Leu Leu Tyr Ile Ser Gln Leu Tyr
1 5 10 15
Met Leu Met Lys Arg Asn Ser Pro His Ile Gly Asp Cys Leu Ser Leu
20 25 30
Leu Phe Leu Gln Glu Lys Lys Gln Lys Glu Val Tyr Thr Leu Leu Ala
35 40 45
Met Met Gln Val Ser Phe Ile Leu Val *
50 55 57

<210> 1828
<211> 102
<212> PRT
<213> Homo sapiens

<400> 1828
Met Gln Pro Ser Gly Leu Glu Gly Pro Gly Thr Phe Gly Arg Trp Pro
1 5 10 15
Leu Leu Ser Leu Leu Leu Leu Leu Leu Leu Leu Gln Pro Val Thr Cys
20 25 30
Ala Tyr Thr Thr Pro Gly Pro Pro Arg Ala Leu Thr Thr Leu Gly Ala
35 40 45
Pro Arg Ala His Thr Met Pro Gly Thr Tyr Ala Pro Ser Thr Thr Leu
50 55 60
Ser Ser Pro Ser Thr Gln Gly Leu Gln Glu Gln Ala Arg Ala Leu Met
65 70 75 80
Arg Asp Phe Pro Leu Val Asp Gly His Asn Asp Leu Pro Leu Val Leu
85 90 95
Arg Gln Val Tyr His Asn
100 102

<210> 1829
<211> 88
<212> PRT
<213> Homo sapiens

<400> 1829
Met Arg Lys Ile Tyr Thr Thr Val Leu Phe Ala Asn Ile Tyr Leu Ala
1 5 10 15
Pro Leu Ser Leu Ile Val Ile Met Tyr Gly Arg Ile Gly Ile Ser Leu
20 25 30
Phe Arg Ala Ala Val Pro His Thr Gly Arg Lys Asn Gln Glu Gln Trp
35 40 45
His Val Val Ser Arg Lys Lys Gln Lys Ile Ile Lys Met Leu Leu Ile
50 55 60
Val Ala Leu Leu Phe Ile Leu Ser Trp Leu Pro Leu Trp Thr Leu Met
65 70 75 80

```
Met Leu Ser Asp Tyr Ala Lys Pro
                85          88


      <210> 1830
      <211> 120
      <212> PRT
      <213> Homo sapiens


      <400> 1830
Met Lys Trp Arg Arg Lys Ser Ala Tyr Trp Lys Ala Leu Lys Val Phe
 1               5                  10                  15
Lys Leu Pro Val Glu Phe Leu Leu Leu Leu Thr Val Pro Val Val Asp
            20                  25                  30
Pro Asp Lys Asp Asp Gln Asn Trp Lys Arg Pro Leu Asn Cys Leu His
        35                  40                  45
Leu Val Ile Ser Pro Leu Val Val Val Leu Thr Leu Gln Ser Gly Thr
    50                  55                  60
Tyr Gly Val Tyr Glu Ile Gly Gly Leu Val Pro Val Trp Val Val Val
65                  70                  75                  80
Val Ile Ala Gly Thr Ala Leu Ala Ser Val Thr Phe Phe Ala Thr Ser
                85                  90                  95
Asp Ser Gln Pro Pro Arg Leu His Trp Leu Phe Ala Phe Leu Gly Phe
            100                 105                 110
Leu Thr Ser Ala Leu Trp Ile Asn
        115                 120


      <210> 1831
      <211> 64
      <212> PRT
      <213> Homo sapiens


      <400> 1831
Met Phe Trp Arg Gly Trp Gly Ala Pro Leu Trp Ala Trp Pro Thr Leu
 1               5                  10                  15
Leu Thr Pro Ile Lys Cys Ser Ser Leu Tyr Asp Ser Phe Phe Ser Pro
            20                  25                  30
Thr Asp Ala Leu Gly Leu Glu Ser Leu Leu Gly Thr Ala Ser Leu Trp
        35                  40                  45
Pro Leu Leu Leu Ser Leu Thr Glu Leu Pro Ala Leu Leu Gln Met  *
    50                  55                  60              63


      <210> 1832
      <211> 89
      <212> PRT
      <213> Homo sapiens


      <400> 1832
Met Gly Ile Lys His Phe Ser Gly Leu Phe Val Leu Leu Cys Ile Gly
 1               5                  10                  15
Phe Gly Leu Ser Ile Leu Thr Thr Ile Gly Glu His Ile Val Tyr Arg
```

```
                20                  25                  30
Leu Leu Leu Pro Arg Ile Lys Asn Lys Ser Lys Leu Gln Tyr Trp Leu
            35                  40                  45
His Thr Ser Gln Arg Leu His Arg Ala Ile Asn Thr Ser Phe Ile Glu
    50                  55                  60
Glu Lys Gln Gln His Phe Lys Thr Lys Arg Val Glu Lys Arg Ser Asn
65                  70                  75                  80
Val Gly Pro Arg Gln Leu Thr Val Trp
                85              89
```

<210> 1833
<211> 60
<212> PRT
<213> Homo sapiens

<400> 1833

```
Met Phe Leu Val Ser Ile Ile Cys Val Thr Leu Phe Phe Pro Ile Val
 1               5                  10                  15
Ala Leu Phe Asp Leu Tyr Ala Thr Leu Ala His Cys Val Tyr Ala Phe
                20                  25                  30
Ser Thr Asp Ser Leu Leu Pro Ala Val Met Leu Thr Ala Leu Pro Arg
            35                  40                  45
Ser Leu Phe Phe Ser Ser Ser Leu Ile Leu Ser Ser
    50                  55                  60
```

<210> 1834
<211> 62
<212> PRT
<213> Homo sapiens

<400> 1834

```
Met Val Pro Ala Ala Gly Ala Leu Leu Trp Val Leu Leu Leu Asn Leu
 1               5                  10                  15
Gly Pro Arg Ala Ala Gly Ala Gln Gly Leu Thr Gln Thr Pro Thr Glu
                20                  25                  30
Met Gln Arg Val Met Leu Arg Phe Gly Cys Ser Val Ile Cys Cys Tyr
            35                  40                  45
Cys Ile Ser Val Arg Thr Gly Arg Ser Arg Glu Thr Gly  *
    50                  55                  60  61
```

<210> 1835
<211> 71
<212> PRT
<213> Homo sapiens

<400> 1835

```
Met Leu Leu Lys Ile Leu Lys Gly Cys Val Val Phe His His Leu Pro
 1               5                  10                  15
Cys Ser Thr Gln Val Tyr Lys Pro Ser Leu Gly Met Trp Gly Phe Leu
                20                  25                  30
```

Ser Pro Leu Trp Glu Val Val Phe Cys His Thr Pro Cys Phe Arg Ala
35 40 45
Gln Pro Gln Leu Asp Arg Ala Gly Ser Ser Phe Leu Ile Tyr Pro Ser
50 55 60
Pro His Ser Thr Ser Asn *
65 70

<210> 1836
<211> 110
<212> PRT
<213> Homo sapiens

<400> 1836
Met Leu Met Tyr Met Phe Tyr Val Leu Pro Phe Cys Gly Leu Ala Ala
1 5 10 15
Tyr Ala Leu Thr Phe Pro Gly Cys Ser Trp Leu Pro Asp Trp Ala Leu
20 25 30
Val Phe Ala Gly Gly Ile Gly Gln Ala Gln Phe Ser His Met Gly Ala
35 40 45
Ser Met His Leu Arg Thr Pro Phe Thr Tyr Arg Val Pro Glu Asp Thr
50 55 60
Trp Gly Cys Phe Phe Val Cys Asn Leu Leu Tyr Ala Leu Gly Pro His
65 70 75 80
Leu Leu Ala Tyr Arg Cys Leu Gln Trp Pro Ala Phe Phe His Gln Pro
85 90 95
Pro Pro Ser Asp Pro Leu Ala Leu His Lys Lys Gln His *
100 105 109

<210> 1837
<211> 91
<212> PRT
<213> Homo sapiens

<400> 1837
Met Leu Leu Leu Leu Thr Trp Pro Tyr Ile Leu Leu Gly Phe Leu Phe
1 5 10 15
Cys Ala Phe Val Val Val Asn Gly Gly Ile Val Ile Gly Asp Arg Ser
20 25 30
Ser His Glu Ala Cys Leu His Phe Pro Gln Leu Phe Tyr Phe Phe Ser
35 40 45
Phe Thr Leu Phe Phe Ser Phe Pro His Leu Leu Ser Pro Ser Lys Ile
50 55 60
Lys Thr Phe Leu Ser Leu Val Trp Lys Arg Arg Ile Leu Phe Phe Val
65 70 75 80
Val Thr Leu Val Ser Val Phe Leu Val Trp Asn
85 90 91

<210> 1838
<211> 201
<212> PRT
<213> Homo sapiens

<400> 1838
Met Pro Ile Gly Leu Arg Gly Leu Met Ile Ala Val Met Leu Ala Ala
1 5 10 15
Leu Met Ser Ser Leu Thr Ser Ile Phe Asn Ser Ser Ser Thr Leu Phe
20 25 30
Thr Met Asp Ile Trp Arg Arg Leu Arg Pro Arg Ser Gly Glu Arg Glu
35 40 45
Leu Leu Leu Val Gly Arg Leu Val Ile Val Ala Leu Ile Gly Val Ser
50 55 60
Val Ala Trp Ile Pro Val Leu Gln Asp Ser Asn Ser Gly Gln Leu Phe
65 70 75 80
Ile Tyr Met Gln Ser Val Thr Ser Ser Leu Ala Pro Pro Val Thr Ala
85 90 95
Val Phe Val Leu Gly Val Phe Trp Arg Arg Ala Asn Glu Gln Gly Ala
100 105 110
Phe Trp Gly Leu Ile Ala Gly Leu Val Val Gly Ala Thr Arg Leu Val
115 120 125
Leu Glu Phe Leu Asn Pro Ala Pro Pro Cys Gly Glu Pro Asp Thr Arg
130 135 140
Pro Ala Val Leu Gly Ser Ile His Tyr Leu His Phe Ala Val Ala Leu
145 150 155 160
Phe Ala Leu Ser Gly Ala Val Val Val Ala Gly Ser Leu Leu Thr Pro
165 170 175
Pro Pro Gln Ser Val Gln Ile Glu Asn Leu Thr Trp Trp Thr Leu Ala
180 185 190
Gln Asp Val Pro Leu Gly Thr Lys Ala
195 200 201

<210> 1839
<211> 130
<212> PRT
<213> Homo sapiens

<221> misc_feature
<222> (1)...(130)
<223> Xaa = any amino acid or nothing

<400> 1839
Met Leu Phe Phe Leu Gln Ser Leu Phe Met Leu Ala Thr Val Val Leu
1 5 10 15
Tyr Phe Ser His Leu Lys Glu Tyr Val Ala Ser Met Val Phe Ser Leu
20 25 30
Ala Leu Gly Trp Thr Asn Met Leu Tyr Tyr Thr Arg Gly Phe Gln Gln
35 40 45
Met Gly Ile Tyr Ala Val Met Ile Glu Lys Met Ile Leu Arg Asp Leu
50 55 60
Cys Arg Phe Met Phe Val Tyr Ile Val Phe Leu Phe Gly Phe Ser Thr
65 70 75 80
Ala Val Val Thr Leu Ile Glu Asp Gly Lys Asn Asp Ser Leu Pro Ser
85 90 95
Glu Ser Thr Ser His Arg Trp Arg Gly Phe Ser Xaa Thr Pro Leu Xaa
100 105 110
Leu Leu His Lys Leu Tyr Ser Thr Cys Leu Glu Leu Ser Asn Ser Thr
115 120 125

Xaa Asp
130

<210> 1840
<211> 47
<212> PRT
<213> Homo sapiens

<400> 1840
Met Asn Arg Val Met Arg Gly Leu Ala Ile Thr Thr Thr Cys Leu Leu
1 5 10 15
Ser Met Leu Gln Ala Ile Thr Ile Ser Pro Ser Ile Leu Trp Asn His
20 25 30
Ala Ala Val Gln Tyr Val His Gly His Ser Leu Val Gln Ala *
35 40 45 46

<210> 1841
<211> 82
<212> PRT
<213> Homo sapiens

<400> 1841
Met Thr Ala Arg Leu Met Arg Ser Leu Leu Ala Ala Gln Leu Thr Phe
1 5 10 15
Val Tyr Arg Val Ala His Leu Met Asn Val Ala Gln Arg Ile Arg Gly
20 25 30
Asn Arg Pro Ile Lys Asn Glu Arg Leu Leu Ala Leu Leu Gly Asp Asn
35 40 45
Glu Lys Met Asn Leu Ser Asp Val Glu Leu Ile Pro Leu Pro Leu Glu
50 55 60
Pro Gln Val Lys Ile Arg Gly Ile Ile Pro Glu Thr Ala Thr Leu Phe
65 70 75 80
Lys Ser
82

<210> 1842
<211> 77
<212> PRT
<213> Homo sapiens

<400> 1842
Met Val Ala Asn Met Phe Tyr Ile Val Val Ile Met Ala Leu Val Leu
1 5 10 15
Leu Ser Phe Gly Val Pro Arg Lys Ala Ile Leu Tyr Pro His Glu Ala
20 25 30
Pro Ser Trp Thr Leu Ala Lys Asp Ile Val Phe His Pro Tyr Trp Met
35 40 45
Ile Phe Gly Glu Val Tyr Ala Tyr Glu Ile Asp Val Cys Ala Asn Asp
50 55 60
Ser Val Ile Pro Gln Ile Cys Gly Pro Ser Thr Arg Pro

65 70 75 77

<210> 1843
<211> 109
<212> PRT
<213> Homo sapiens

<400> 1843
Met Met His Asn Ile Ile Val Lys Glu Leu Ile Val Thr Phe Phe Leu
1 5 10 15
Gly Ile Thr Val Val Gln Met Leu Ile Ser Val Thr Gly Leu Lys Gly
20 25 30
Val Glu Ala Gln Asn Gly Ser Glu Ser Glu Val Phe Val Gly Lys Tyr
35 40 45
Glu Thr Leu Val Phe Tyr Trp Pro Ser Leu Leu Cys Leu Ala Phe Leu
50 55 60
Leu Gly Arg Phe Leu His Met Phe Val Lys Ala Leu Arg Val His Leu
65 70 75 80
Gly Trp Glu Leu Gln Val Glu Glu Lys Ser Val Leu Glu Val His Gln
85 90 95
Gly Glu His Val Lys Gln Leu Leu Arg Ile Pro Arg Pro
100 105 109

<210> 1844
<211> 85
<212> PRT
<213> Homo sapiens

<221> misc_feature
<222> (1)...(85)
<223> Xaa = any amino acid or nothing

<400> 1844
Met Thr Ile His Leu Cys Ser Asn Leu Met Cys His Phe Leu Gln Arg
1 5 10 15
Met Gly Thr Ile Leu Leu Cys Pro Asn Met Gln Pro His Gln Asn Leu
20 25 30
Thr Thr Val Ile Cys Ser Lys Gly Asn Leu Leu Arg Ala Val Lys Gly
35 40 45
Ser Lys Ser Leu Arg Asn Ala Arg Lys Tyr Pro Phe His His Pro Pro
50 55 60
Xaa Xaa Glu Pro Pro Asn Gly Gly Gln Thr Arg Xaa Gly Gly Ala Arg
65 70 75 80
Phe Lys Gln Pro Thr
85

<210> 1845
<211> 110
<212> PRT
<213> Homo sapiens

<400> 1845

```
Met Tyr Ala Leu Tyr Ile Thr Val His Gly Tyr Phe Leu Ile Thr Phe
 1               5                  10                  15
Leu Phe Gly Met Val Val Leu Ala Leu Val Val Trp Lys Ile Phe Thr
            20                  25                  30
Leu Ser Arg Ala Thr Ala Val Lys Glu Arg Gly Lys Asn Arg Lys Lys
        35                  40                  45
Val Leu Thr Leu Leu Gly Leu Ser Ser Leu Val Gly Val Thr Trp Gly
    50                  55                  60
Leu Ala Ile Phe Thr Pro Leu Gly Leu Ser Thr Val Tyr Ile Phe Ala
 65                 70                  75                  80
Leu Phe Asn Ser Leu Gln Gly Val Phe Ile Cys Cys Trp Phe Thr Ile
                85                  90                  95
Leu Tyr Leu Pro Ser Gln Ser Thr Thr Val Ser Ser Ser Thr
            100                 105                 110
```

<210> 1846
<211> 94
<212> PRT
<213> Homo sapiens

<400> 1846

```
Met Thr Glu Pro Pro Gly Ala Ser Ser His Leu Arg Gln Ala Leu Arg
 1               5                  10                  15
Cys Cys Gln Trp Leu Ala Gly Ile Pro Ser Gln Trp Val Leu Phe Trp
            20                  25                  30
Glu Val Leu Trp Lys Trp Val Leu Gln Thr Asp Ala Ala Trp Ser Pro
        35                  40                  45
Gly Phe Ser Pro Leu Pro Arg Gly Met Tyr Gln His Pro Ala Leu Pro
    50                  55                  60
Glu Met Pro Ser Pro Phe Leu Gly Ile Leu Arg Leu Glu Tyr Val Lys
 65                 70                  75                  80
Leu Leu Gly Leu Cys Met Cys Leu Ser Thr Gly Ser Ser  *
                85                  90          93
```

<210> 1847
<211> 1300
<212> PRT
<213> Homo sapiens

<400> 1847

```
Met Ala Trp Lys Thr Leu Pro Ile Tyr Leu Leu Leu Leu Leu Ser Val
 1               5                  10                  15
Phe Val Ile Gln Gln Val Ser Ser Gln Asp Leu Ser Ser Cys Ala Gly
            20                  25                  30
Arg Cys Gly Glu Gly Tyr Ser Arg Asp Ala Thr Cys Asn Cys Asp Tyr
        35                  40                  45
Asn Cys Gln His Tyr Met Glu Cys Cys Pro Asp Phe Lys Arg Val Cys
    50                  55                  60
Thr Ala Glu Leu Ser Cys Lys Gly Arg Cys Phe Glu Ser Phe Glu Arg
 65                 70                  75                  80
Gly Arg Glu Cys Asp Cys Asp Ala Gln Cys Lys Lys Tyr Asp Lys Cys
```

85 90 95
Cys Pro Asp Tyr Glu Ser Phe Cys Ala Glu Val His Asn Pro Thr Ser
100 105 110
Pro Pro Ser Ser Lys Lys Ala Pro Pro Pro Ser Gly Ala Ser Gln Thr
115 120 125
Ile Lys Ser Thr Thr Lys Arg Ser Pro Lys Pro Pro Asn Lys Lys Lys
130 135 140
Thr Lys Lys Val Ile Glu Ser Glu Glu Ile Thr Glu Glu His Ser Val
145 150 155 160
Ser Glu Asn Gln Glu Ser Ser Ser Ser Ser Ser Ser Ser Ser Ser Ser
165 170 175
Ser Thr Ile Trp Lys Ile Lys Ser Ser Lys Asn Ser Ala Ala Asn Arg
180 185 190
Glu Leu Gln Lys Lys Leu Lys Val Lys Asp Asn Lys Lys Asn Arg Thr
195 200 205
Lys Lys Lys Pro Thr Pro Lys Pro Pro Val Val Asp Glu Ala Gly Ser
210 215 220
Gly Leu Asp Asn Gly Asp Phe Lys Val Thr Thr Pro Asp Thr Ser Thr
225 230 235 240
Thr Gln His Asn Lys Val Ser Thr Ser Pro Lys Ile Thr Thr Ala Lys
245 250 255
Pro Ile Asn Pro Arg Pro Ser Leu Pro Pro Asn Ser Asp Thr Ser Lys
260 265 270
Glu Thr Ser Leu Thr Val Asn Lys Glu Thr Thr Val Glu Thr Lys Glu
275 280 285
Thr Thr Thr Thr Asn Lys Gln Thr Ser Thr Asp Gly Lys Glu Lys Thr
290 295 300
Thr Ser Ala Lys Glu Thr Gln Ser Ile Glu Lys Thr Ser Ala Lys Asp
305 310 315 320
Leu Ala Pro Thr Ser Lys Val Leu Ala Lys Pro Thr Pro Lys Ala Glu
325 330 335
Thr Thr Thr Lys Gly Pro Ala Leu Thr Thr Pro Lys Glu Pro Thr Pro
340 345 350
Thr Thr Pro Lys Glu Pro Ala Ser Thr Thr Pro Lys Glu Pro Thr Pro
355 360 365
Thr Thr Ile Lys Ser Ala Pro Thr Thr Pro Lys Glu Pro Ala Pro Thr
370 375 380
Thr Thr Lys Ser Ala Pro Thr Thr Pro Lys Glu Pro Ala Pro Thr Thr
385 390 395 400
Thr Lys Glu Pro Ala Pro Thr Thr Pro Lys Glu Pro Ala Pro Thr Thr
405 410 415
Thr Lys Glu Pro Ala Pro Thr Thr Thr Lys Ser Ala Pro Thr Thr Pro
420 425 430
Lys Glu Pro Ala Pro Thr Thr Pro Lys Lys Pro Ala Pro Thr Thr Pro
435 440 445
Lys Glu Pro Ala Pro Thr Thr Pro Lys Glu Pro Thr Pro Thr Thr Pro
450 455 460
Lys Glu Pro Ala Pro Thr Thr Lys Glu Pro Ala Pro Thr Thr Pro Lys
465 470 475 480
Glu Pro Ala Pro Thr Ala Pro Lys Lys Pro Ala Pro Thr Thr Pro Lys
485 490 495
Glu Pro Ala Pro Thr Thr Pro Lys Glu Pro Ala Pro Thr Thr Thr Lys
500 505 510
Glu Pro Ser Pro Thr Thr Pro Lys Glu Pro Ala Pro Thr Thr Thr Lys
515 520 525
Ser Ala Pro Thr Thr Thr Lys Glu Pro Ala Pro Thr Thr Thr Lys Ser
530 535 540
Ala Pro Thr Thr Pro Lys Glu Pro Ser Pro Thr Thr Thr Lys Glu Pro
545 550 555 560

Ala Pro Thr Thr Pro Lys Glu Pro Ala Pro Thr Thr Pro Lys Lys Pro
565 570 575
Ala Pro Thr Thr Pro Lys Glu Pro Ala Pro Thr Thr Pro Lys Glu Pro
580 585 590
Ala Pro Thr Thr Thr Lys Lys Pro Ala Pro Thr Ala Pro Lys Glu Pro
595 600 605
Ala Pro Thr Thr Pro Lys Glu Thr Ala Pro Thr Thr Pro Lys Lys Leu
610 615 620
Thr Pro Thr Thr Pro Glu Lys Leu Ala Pro Thr Thr Pro Glu Lys Pro
625 630 635 640
Ala Pro Thr Thr Pro Glu Glu Leu Ala Pro Thr Thr Pro Glu Glu Pro
645 650 655
Thr Pro Thr Thr Pro Glu Glu Pro Ala Pro Thr Thr Pro Lys Ala Ala
660 665 670
Ala Pro Asn Thr Pro Lys Glu Pro Ala Pro Thr Thr Pro Lys Glu Pro
675 680 685
Ala Pro Thr Thr Pro Lys Glu Pro Ala Pro Thr Thr Pro Lys Glu Thr
690 695 700
Ala Pro Thr Thr Pro Lys Gly Thr Ala Pro Thr Thr Leu Lys Glu Pro
705 710 715 720
Ala Pro Thr Thr Pro Lys Lys Pro Ala Pro Lys Glu Leu Ala Pro Thr
725 730 735
Thr Thr Lys Glu Pro Thr Ser Thr Thr Ser Asp Lys Pro Ala Pro Thr
740 745 750
Thr Pro Lys Gly Thr Ala Pro Thr Thr Pro Lys Glu Pro Ala Pro Thr
755 760 765
Thr Pro Lys Glu Pro Ala Pro Thr Thr Pro Lys Gly Thr Ala Pro Thr
770 775 780
Thr Leu Lys Glu Pro Ala Pro Thr Thr Pro Lys Lys Pro Ala Pro Lys
785 790 795 800
Glu Leu Ala Pro Thr Thr Thr Lys Gly Pro Thr Ser Thr Thr Ser Asp
805 810 815
Lys Pro Ala Pro Thr Thr Pro Lys Glu Thr Ala Pro Thr Thr Pro Lys
820 825 830
Glu Pro Ala Pro Thr Thr Pro Lys Lys Pro Ala Pro Thr Thr Pro Glu
835 840 845
Thr Pro Pro Pro Thr Thr Ser Glu Val Ser Thr Pro Thr Thr Thr Lys
850 855 860
Glu Pro Thr Thr Ile His Lys Ser Pro Asp Glu Ser Thr Pro Glu Leu
865 870 875 880
Ser Ala Glu Pro Thr Pro Lys Ala Leu Glu Asn Ser Pro Lys Glu Pro
885 890 895
Gly Val Pro Thr Thr Lys Thr Pro Ala Ala Thr Lys Pro Glu Met Thr
900 905 910
Thr Thr Ala Lys Asp Lys Thr Thr Glu Arg Asp Leu Arg Thr Thr Pro
915 920 925
Glu Thr Thr Thr Ala Ala Pro Lys Met Thr Lys Glu Thr Ala Thr Thr
930 935 940
Thr Glu Lys Thr Thr Glu Ser Lys Ile Thr Ala Thr Thr Thr Gln Val
945 950 955 960
Thr Ser Thr Thr Thr Gln Asp Thr Thr Pro Phe Lys Ile Thr Thr Leu
965 970 975
Lys Thr Thr Thr Leu Ala Pro Lys Val Thr Thr Thr Lys Lys Thr Ile
980 985 990
Thr Thr Thr Glu Ile Met Asn Lys Pro Glu Glu Thr Ala Lys Pro Lys
995 1000 1005
Asp Arg Ala Thr Asn Ser Lys Ala Thr Thr Pro Lys Pro Gln Lys Pro
1010 1015 1020
Thr Lys Ala Pro Lys Lys Pro Thr Ser Thr Lys Lys Pro Lys Thr Met

1025 1030 1035 1040
Pro Arg Val Arg Lys Pro Lys Thr Thr Pro Thr Pro Arg Lys Met Thr
1045 1050 1055
Ser Thr Met Pro Glu Leu Asn Pro Thr Ser Arg Ile Ala Glu Ala Met
1060 1065 1070
Leu Gln Thr Thr Thr Arg Pro Asn Gln Thr Pro Asn Ser Lys Leu Val
1075 1080 1085
Glu Val Asn Pro Lys Ser Glu Asp Ala Gly Gly Ala Glu Gly Glu Thr
1090 1095 1100
Pro His Met Leu Leu Arg Pro His Val Phe Met Pro Glu Val Thr Pro
1105 1110 1115 1120
Asp Met Asp Tyr Leu Pro Arg Val Pro Asn Gln Gly Ile Ile Ile Asn
1125 1130 1135
Pro Met Leu Ser Asp Glu Thr Asn Ile Cys Asn Gly Lys Pro Val Asp
1140 1145 1150
Gly Leu Thr Thr Leu Arg Asn Gly Thr Leu Val Ala Phe Arg Gly His
1155 1160 1165
Tyr Phe Trp Met Leu Ser Pro Phe Ser Pro Pro Ser Pro Ala Arg Arg
1170 1175 1180
Ile Thr Glu Val Trp Gly Ile Pro Ser Pro Ile Asp Thr Val Phe Thr
1185 1190 1195 1200
Arg Cys Asn Cys Glu Gly Lys Thr Phe Phe Phe Lys Asp Ser Gln Tyr
1205 1210 1215
Trp Arg Phe Thr Asn Asp Ile Lys Asp Ala Gly Tyr Pro Lys Pro Ile
1220 1225 1230
Phe Lys Gly Phe Gly Gly Leu Thr Gly Gln Ile Val Ala Ala Leu Ser
1235 1240 1245
Thr Ala Lys Tyr Lys Asn Trp Pro Glu Ser Val Tyr Phe Phe Lys Arg
1250 1255 1260
Gly Gly Ser Ile Gln Gln Tyr Ile Tyr Lys Gln Glu Pro Val Gln Lys
1265 1270 1275 1280
Cys Pro Gly Arg Arg Pro Ala Leu Asn Tyr Pro Val Tyr Gly Glu Thr
1285 1290 1295
Asp Thr Gly *
1299

<210> 1848
<211> 103
<212> PRT
<213> Homo sapiens

<400> 1848
Met Asn Pro Ala Val Arg Gln Arg Cys Leu Leu Phe Cys Phe Gln Gln
1 5 10 15
Lys Leu Ile Leu Ser His Phe Phe Leu Leu Gln Val Pro Gln Trp Cys
20 25 30
Ala Glu Tyr Cys Leu Ser Ile His Tyr Gln His Gly Gly Val Ile Cys
35 40 45
Thr Gln Val His Lys Gln Thr Val Val Gln Leu Ala Leu Arg Val Ala
50 55 60
Asp Glu Met Asp Val Asn Ile Gly His Glu Val Gly Tyr Val Ile Pro
65 70 75 80
Phe Glu Asn Cys Cys Thr Asn Glu Thr Ile Leu Arg Leu Val Cys Gly
85 90 95
Val Gln Ser Ala Pro Cys *
100 102

<210> 1849
<211> 50
<212> PRT
<213> Homo sapiens

<400> 1849
Met Ser Arg Phe Leu Leu Pro Arg Glu Gly Cys Leu Leu Ile Val Phe
1 5 10 15
Met Leu Cys Glu Lys Thr Leu Pro Phe Leu Phe Thr Leu Lys Glu Tyr
20 25 30
Thr Phe Ile Pro Glu His Arg Thr Thr Asp Ile Asn Cys Val Asn Thr
35 40 45
His Glu
50

<210> 1850
<211> 84
<212> PRT
<213> Homo sapiens

<400> 1850
Met Arg Leu His Ser Lys Gly Ser Gln Asp Pro Ser Thr Lys Val His
1 5 10 15
Ile Lys Ala Leu Gln Thr Val Thr Ser Phe Leu Met Leu Phe Ala Ile
20 25 30
Tyr Phe Leu Cys Ile Ile Thr Ser Thr Trp Asn Leu Arg Thr Gln Gln
35 40 45
Ser Lys Leu Val Leu Leu Leu Cys Gln Thr Val Ala Ile Met Tyr Pro
50 55 60
Ser Phe His Ser Phe Ile Leu Ile Met Gly Ser Arg Lys Leu Lys Gln
65 70 75 80
Thr Phe Leu Ser
84

<210> 1851
<211> 51
<212> PRT
<213> Homo sapiens

<400> 1851
Met Ala Ala Cys Lys Leu Leu Lys His Leu Asn Gly Phe Ser Leu Leu
1 5 10 15
Leu Pro Arg Leu Glu Cys Asn Gly Val Ile Ser Val His Cys Asn Pro
20 25 30
Leu Pro Pro Gly Phe Lys Arg Phe Ser Cys Pro Ser Leu Leu Ser Ser
35 40 45
Trp Asp *
50

<210> 1852
<211> 54
<212> PRT
<213> Homo sapiens

<400> 1852
Met Lys Thr Lys Cys Lys Pro Asn Ile Thr Phe Phe Asn Thr Ile Ile
1 5 10 15
Cys Phe Phe Leu Thr Phe Leu Phe Cys Ile Tyr Ile Asp Ser Leu Leu
20 25 30
Cys Thr Val Pro Lys Asn Pro Ala Gln Ala Val Gln Leu Asn Arg Asp
35 40 45
His Thr Lys Val His *
50 53

<210> 1853
<211> 129
<212> PRT
<213> Homo sapiens

<400> 1853
Met Ala Val Val Arg Val Met Val Val Val Arg Val Thr Ala Val Val
1 5 10 15
Arg Val Met Val Val Val Arg Val Val Val Val Arg Val Met Val Val
20 25 30
Val Arg Ile Thr Ala Val Leu Arg Val Met Val Val Val Arg Ile Met
35 40 45
Ala Val Ile Arg Val Met Val Val Val Arg Val Thr Ala Ile Val Gly
50 55 60
Val Met Val Val Ile Arg Val Thr Ala Ile Val Ser Ile Met Val Val
65 70 75 80
Val Arg Val Met Val Val Val Arg Val Met Val Val Ala Arg Pro Met
85 90 95
Val Val Val Arg Val Met Ala Val Val Arg Val Met Ala Asp Ser Ala
100 105 110
Leu Arg Ala Ile Cys Ser Ser Ser Leu Asn Val Thr Phe Ser Leu Glu
115 120 125 128
*

<210> 1854
<211> 190
<212> PRT
<213> Homo sapiens

<221> misc_feature
<222> (1)...(190)
<223> Xaa = any amino acid or nothing

<400> 1854

```
Met Ser Cys Phe Gly Leu Leu Leu Gly Gly Leu Thr Pro Arg Val Leu
 1               5                  10                  15
Ser Thr Glu Glu Gln Leu Pro Pro Gly Phe Pro Ser Ile Asp Met Gly
            20                  25                  30
Pro Gln Leu Lys Val Val Glu Lys Ala Arg Thr Ala Thr Met Leu Cys
        35                  40                  45
Ala Ala Gly Gly Asn Pro Asp Pro Glu Ile Ser Trp Phe Lys Asp Phe
    50                  55                  60
Leu Pro Val Asp Pro Ala Thr Ser Asn Gly Arg Ile Lys Gln Leu Arg
65                  70                  75                  80
Ser Gly Glu Gln Arg Ala Gly Val Lys Gly Pro Cys Arg Pro Gln Asn
                85                  90                  95
Lys Arg Leu Val Arg Ser Gln His Ser Leu Leu Pro Trp Ala Trp Ala
            100                 105                 110
Pro Pro Gly Leu Ser Gly Gly Tyr Leu Val Gly Trp Ala Gly Ser Tyr
        115                 120                 125
Cys Arg Cys Ala Trp Leu Arg Glu Glu Ser Ser Trp Leu Ala Val Pro
    130                 135                 140
Leu Pro Ser Ser Asp Cys Gln Thr Pro Asp Phe Gly Pro Val Leu Pro
145                 150                 155                 160
Leu Pro Ala His Val Met Cys Gln Cys Gly Gly Leu Phe Lys Gly Ala
                165                 170                 175
Leu Trp Met Leu Thr Leu Leu Leu Pro Cys Xaa Leu Ala  *
            180                 185                 189


     &lt;210&gt; 1855
     &lt;211&gt; 78
     &lt;212&gt; PRT
     &lt;213&gt; Homo sapiens


     &lt;400&gt; 1855
Met Val Val Ser Ala Trp Ile Gly Leu Glu Ala Thr Val Val Ala Ala
 1               5                  10                  15
Cys Leu Ala Leu Leu Gly Ser Val Val Arg Glu Thr Ser Thr Ser Ala
            20                  25                  30
Ser Pro Thr Pro Ala Ala Leu Arg Ala Ala Trp Thr Val Tyr Ser Ser
        35                  40                  45
Pro Met Thr Thr Cys Val Phe Ala Val Val Pro Leu Leu Ala Gly Thr
    50                  55                  60
Val Lys Pro Ser Ser Met Cys Val Pro Arg Cys Pro Ala  *
65                  70                  75      77


     &lt;210&gt; 1856
     &lt;211&gt; 67
     &lt;212&gt; PRT
     &lt;213&gt; Homo sapiens


     &lt;400&gt; 1856
Met Thr Asn Trp Met Leu Leu Leu Ala Ser Arg Ile Phe Gln Ser Leu
 1               5                  10                  15
Ala Ile Pro Lys Gln Leu Gly Leu Arg Arg Glu Met Pro Ser Gly Ser
            20                  25                  30
Pro Thr Thr Asn Ser Ser Ser Gly Cys Ile Arg Asn Leu Glu Tyr Ser
```

```
        35                  40                  45
Thr Leu Met Gly Ser Glu Met Pro Met Ala Leu Ala Ala Glu Thr Trp
    50                  55                  60
Leu Leu  *
 65  66
```

<210> 1857
<211> 107
<212> PRT
<213> Homo sapiens

<400> 1857

```
Met Leu Leu Met Phe Leu Leu Ala Thr Cys Leu Leu Ala Ile Ile Phe
  1               5                  10                  15
Val Pro Gln Glu Met Gln Thr Leu Arg Val Val Leu Ala Thr Leu Gly
             20                  25                  30
Val Gly Ala Ala Ser Leu Gly Ile Thr Cys Ser Thr Ala Gln Glu Asn
        35                  40                  45
Glu Leu Ile Pro Ser Ile Ile Arg Gly Arg Ala Thr Gly Ile Thr Gly
    50                  55                  60
Asn Phe Ala Asn Ile Gly Gly Ala Leu Ala Ser Leu Val Met Ile Leu
 65                  70                  75                  80
Ser Ile Tyr Ser Arg Pro Leu Pro Trp Ile Ile Tyr Gly Val Phe Ala
                 85                  90                  95
Ile Leu Ser Gly Leu Val Val Leu Leu Leu Pro
            100                 105     107
```

<210> 1858
<211> 134
<212> PRT
<213> Homo sapiens

<400> 1858

```
Met Ile Pro Pro Ala Ile Phe Trp Val Leu Ile Ile Phe Gly Trp Thr
  1               5                  10                  15
Leu Val Tyr Gly Phe Val Tyr Phe Thr Thr Gly Glu Thr Ile Met Asp
             20                  25                  30
Lys Leu Leu Arg Val Leu Tyr Trp Ile Leu Val Lys Thr Phe Phe Arg
        35                  40                  45
Glu Ile Ser Val Ser His Gln Glu Arg Ile Pro Lys Asp Lys Pro Val
    50                  55                  60
Met Leu Val Cys Ala Pro His Ala Asn Gln Phe Val Asp Gly Met Val
 65                  70                  75                  80
Ile Ser Thr His Leu Asp Arg Lys Val Tyr Phe Val Gly Ala Ala Ser
                 85                  90                  95
Ser Phe Arg Lys Tyr Lys Val Val Gly Leu Phe Met Lys Leu Met Ala
            100                 105                 110
Ser Ile Ile Ser Gly Glu Arg His Gln Asp Val Lys Lys Val Leu Thr
        115                 120                 125
Gly Met Ala Thr Glu Lys
    130             134
```

<210> 1859
<211> 82
<212> PRT
<213> Homo sapiens

<400> 1859
Met Phe Tyr Val Lys Ala Glu Phe Leu Val Ser Phe Ser Cys Pro Trp
1 5 10 15
Leu Thr Ala Cys Ala Leu Leu Met Ser Cys Ser Trp Phe Leu Thr Leu
20 25 30
Thr Ile Leu Ser Val Lys Gly Gly Thr Pro Ala Gly Met Leu Asp Gln
35 40 45
Lys Lys Gly Lys Phe Ala Trp Phe Ser His Ser Thr Glu Thr His Gly
50 55 60
Asn Val Pro Leu Cys Ser Val Cys Val Asn Ala Cys Gly Cys Ile Pro
65 70 75 80
Asp *
81

<210> 1860
<211> 46
<212> PRT
<213> Homo sapiens

<400> 1860
Met Pro Leu Ser Pro Leu Leu Phe His Leu Gly Pro Phe Pro Phe Lys
1 5 10 15
Ala Glu Ser Trp Leu Asn Phe Leu Pro Pro Pro Phe Phe Pro Leu Leu
20 25 30
Pro Leu Leu Phe Leu Ala Lys Ala Glu Ile Gln Trp Ala *
35 40 45

<210> 1861
<211> 128
<212> PRT
<213> Homo sapiens

<400> 1861
Met Thr Ile Phe Phe Ser Leu Leu Val Leu Ala Ile Cys Ile Ile Leu
1 5 10 15
Val His Leu Leu Ile Arg Tyr Arg Leu His Phe Leu Pro Glu Ser Val
20 25 30
Ala Val Val Ser Leu Gly Ile Leu Met Gly Ala Val Ile Lys Ile Ile
35 40 45
Glu Phe Lys Lys Leu Ala Asn Trp Lys Glu Glu Glu Met Phe Arg Pro
50 55 60
Asn Met Phe Phe Leu Leu Leu Leu Pro Pro Ile Ile Phe Glu Ser Gly
65 70 75 80
Tyr Ser Leu His Lys Gly Asn Phe Phe Gln Asn Ile Gly Ser Ile Thr
85 90 95
Leu Phe Ala Val Phe Gly Thr Ala Ile Ser Ala Phe Val Val Gly Gly

```
            100                 105                 110
Gly Ile Tyr Phe Leu Gly Gln Ala His Val Ile Ser Lys Leu Asn Met
        115                 120                 125             128
```

<210> 1862
<211> 58
<212> PRT
<213> Homo sapiens

<400> 1862

```
Met Trp Asp Met Leu Pro Trp Gly Ile Thr Trp Val Leu Leu Thr Thr
 1               5                  10                  15
Gln Leu His Ser Pro Leu Leu Tyr Val Ile Gly Phe Thr Tyr Trp Val
            20                  25                  30
Cys Lys Gly Asp Arg Asp Ser Tyr Leu Glu Glu Asn Ser Arg Glu Thr
        35                  40                  45
Ala Ser Val Tyr Thr Ser Val Leu Ser  *
    50                  55      57
```

<210> 1863
<211> 50
<212> PRT
<213> Homo sapiens

<400> 1863

```
Met Thr Gln Asp Leu Val Leu Thr Val Pro Phe Met Gly Cys Leu Leu
 1               5                  10                  15
Ile Leu Val Asp Gly Leu Lys Pro Asn Arg Pro Ala Tyr Ile Gln Thr
            20                  25                  30
Gly Ser Gln Ala Thr Gln Ala Gly Val Gln Trp His Asn Tyr Gly Ser
        35                  40                  45
Leu  *
 49
```

<210> 1864
<211> 90
<212> PRT
<213> Homo sapiens

<400> 1864

```
Met Val Ala Ser Ala Ala Gln Leu Leu Ser His Val Cys Leu Gly Gly
 1               5                  10                  15
Leu Gln Leu Leu His Ser Phe Leu Ser Ser Leu Gln Leu Pro Ala Leu
            20                  25                  30
Leu Leu Lys Leu Ala Pro Glu Ala Leu Ala Leu Phe Thr Ser Ile Leu
        35                  40                  45
Lys Ser Ala Leu Val Val His Asp Phe Ser Thr Gln Leu Glu Leu Glu
    50                  55                  60
```

Gly Val Glu Leu Leu Val Cys Ser Pro Leu Glu Ala Leu Gly Pro Leu
65 70 75 80
Leu Cys Leu Gly Glu Leu Gly Leu Gln Ala
85 90

<210> 1865
<211> 125
<212> PRT
<213> Homo sapiens

<400> 1865
Met Arg Leu Gly Leu Leu Leu Leu Ala Arg His Trp Cys Ile Ala Gly
1 5 10 15
Val Phe Pro Gln Lys Phe Asp Gly Asp Ser Ala Tyr Val Gly Met Ser
20 25 30
Asp Gly Asn Pro Glu Leu Leu Ser Thr Ser Gln Thr Tyr Asn Gly Gln
35 40 45
Ser Glu Asn Asn Glu Asp Tyr Glu Ile Pro Pro Ile Thr Pro Pro Asn
50 55 60
Leu Pro Glu Pro Ser Leu Leu His Leu Gly Asp His Glu Ala Ser Tyr
65 70 75 80
His Ser Leu Cys His Gly Leu Thr Pro Asn Gly Leu Leu Pro Ala Tyr
85 90 95
Ser Tyr Gln Ala Met Asp Leu Pro Ala Ile Met Val Ser Asn Met Leu
100 105 110
Ala Gln Asp Ser His Leu Leu Ser Gly Gln Leu Pro Thr
115 120 125

<210> 1866
<211> 129
<212> PRT
<213> Homo sapiens

<400> 1866
Met Cys Phe Leu Asn Lys Leu Leu Leu Leu Ala Ala Leu Asp Trp Leu
1 5 10 15
Phe Gln Ile Pro Thr Val Pro Glu Asp Leu Phe Phe Leu Glu Glu Gly
20 25 30
Pro Ser Tyr Ala Phe Glu Val Asp Thr Val Ala Pro Glu His Gly Leu
35 40 45
Asp Asn Ala Pro Val Val Asp Gln Gln Leu Leu Tyr Thr Cys Cys Pro
50 55 60
Tyr Ile Gly Glu Leu Arg Lys Leu Leu Ala Ser Trp Val Ser Gly Ser
65 70 75 80
Ser Gly Arg Ser Gly Gly Phe Met Arg Lys Ile Thr Pro Thr Thr Thr
85 90 95
Thr Ser Leu Gly Ala Gln Pro Ser Gln Thr Ser Gln Gly Leu Gln Ala
100 105 110
Gln Leu Ala Gln Ala Phe Phe His Asn Gln Pro Pro Ser Leu Arg Arg
115 120 125
Thr
129

<210> 1867
<211> 80
<212> PRT
<213> Homo sapiens

<400> 1867
Met Met Arg Leu Glu Lys Phe Val Thr Trp Ser Val Met Ala Leu Gly
1 5 10 15
Trp Phe Val Phe Arg Gln Gln Asn Cys Trp Ala Leu Trp Ser Lys Ser
20 25 30
Val Leu Ile Ser Trp Ser Arg Pro Leu Thr Arg Ser Met Ser Asp Leu
35 40 45
Arg Arg Lys Arg Thr Ala His Glu Arg Ala Lys Glu Leu Tyr Ser Ser
50 55 60
Gly Glu Phe Ser Ser Gly Arg Lys Trp Gly Asp Asp Ala Pro Lys Glu
65 70 75 80

<210> 1868
<211> 113
<212> PRT
<213> Homo sapiens

<400> 1868
Met Leu Val Trp Leu Tyr Gly Thr Ile Arg Trp Pro Ala Leu Gly Ala
1 5 10 15
Pro Arg Trp Trp Pro Trp Val Trp Pro Pro Gly Val Trp Ser Gly Ile
20 25 30
Glu Thr Pro Ser Ser Thr Pro Arg Ala Arg Ser Leu Arg Gly Thr Gly
35 40 45
Gly Ala Val Thr Arg Arg Thr Gly Ser Ser Phe Pro Trp Thr Thr Thr
50 55 60
Thr Arg Pro Ser Ser Trp Trp Thr Thr Ala His Thr Ala Ala Trp Gly
65 70 75 80
Ala Arg Thr Ala Ser Ala Cys Ala Trp Ser Pro Thr Ser His Ser Lys
85 90 95
Thr Arg Pro Trp Gln Gly Leu Glu Leu Thr Ser Leu Ala Cys Ser Ser
100 105 110 112
*

<210> 1869
<211> 72
<212> PRT
<213> Homo sapiens

<400> 1869
Met Phe Leu Trp Val Lys Arg Leu Leu Phe Ala Ala Ser Leu Leu Ala
1 5 10 15

```
Ser Asp Ser Ser Thr Ile Leu Cys Ser Arg Asp Leu Ile Leu Glu Ser
            20                  25                  30
Ile Ala Leu Ile Ile Ala Phe Cys Ser Leu Arg Ile Leu Pro Phe Ser
        35                  40                  45
Trp Ala Ser Ser Ser Cys Leu Cys Ile Met Phe Ser Ser Val Ser Leu
    50                  55                  60
Ser Ala Arg Ser Phe Phe Ile  *
65                  70  71
```

<210> 1870
<211> 197
<212> PRT
<213> Homo sapiens

<400> 1870

```
Met Arg Thr Leu Leu Thr Ile Leu Thr Val Gly Ser Leu Ala Ala His
 1               5                  10                  15
Ala Pro Glu Asp Pro Ser Asp Leu Leu Gln His Val Lys Phe Gln Ser
            20                  25                  30
Ser Asn Phe Glu Asn Ile Leu Thr Trp Asp Ser Gly Pro Glu Gly Thr
        35                  40                  45
Pro Asp Thr Val Tyr Ser Ile Glu Tyr Lys Thr Tyr Gly Glu Arg Asp
    50                  55                  60
Trp Val Ala Lys Lys Gly Cys Gln Arg Ile Thr Arg Lys Ser Cys Asn
65                  70                  75                  80
Leu Thr Val Glu Thr Gly Asn Leu Thr Glu Leu Tyr Tyr Ala Arg Val
                85                  90                  95
Thr Ala Val Ser Ala Gly Gly Arg Ser Ala Thr Lys Met Thr Asp Arg
            100                 105                 110
Phe Ser Ser Leu Gln His Thr Thr Leu Lys Pro Pro Asp Val Thr Cys
        115                 120                 125
Ile Ser Lys Val Arg Ser Ile Gln Met Ile Val His Pro Thr Pro Thr
    130                 135                 140
Pro Ile Arg Ala Gly Asp Gly His Arg Leu Thr Leu Glu Asp Ile Phe
145                 150                 155                 160
His Asp Leu Phe Tyr His Leu Glu Leu Gln Val Asn Arg Thr Tyr Gln
                165                 170                 175
Met Val Ser Val Cys Cys Thr Leu Val Phe Leu Cys Leu Gly Ser Leu
            180                 185                 190
Phe Pro Pro Asn  *
        195 196
```

<210> 1871
<211> 75
<212> PRT
<213> Homo sapiens

<400> 1871

```
Met Glu Tyr Arg Leu Gln Lys Gly Ala Gly Phe His Leu Asp Leu Phe
 1               5                  10                  15
Cys Val Ala Val Leu Met Leu Leu Thr Ser Ala Leu Gly Leu Pro Trp
            20                  25                  30
Tyr Val Ser Ala Thr Val Ile Ser Leu Ala His Met Asp Ser Leu Arg
```

35 40 45
Arg Glu Ser Arg Ala Cys Ala Pro Gly Glu Arg Pro Asn Phe Leu Gly
50 55 60
Ile Arg Glu Gln Arg Leu Thr Gly Leu Val Val
65 70 75

<210> 1872
<211> 84
<212> PRT
<213> Homo sapiens

<400> 1872
Met Pro Phe Ser Thr Cys Thr Ala Leu Pro Ser Trp Ala Thr Leu Ser
1 5 10 15
Thr Trp Ser Trp Thr Pro Lys Val Ser Leu Ala Gly Glu Glu Arg Gly
20 25 30
Glu Thr Cys Gln Pro Asp Pro Phe Pro Pro His Pro Ser Cys Ser Val
35 40 45
Gly Arg Thr Pro Pro His Ser Ser Leu Gly Ser Pro Pro Thr Thr Leu
50 55 60
Phe Leu Ser Pro Leu Leu Arg Val Glu Ser Arg Gly Ala Lys Cys Val
65 70 75 80
Val Cys Cys *
83

<210> 1873
<211> 51
<212> PRT
<213> Homo sapiens

<400> 1873
Met Cys Gly Ser Pro Glu Arg Leu Cys Val Arg Cys Ala Arg Val Cys
1 5 10 15
Ala Val Phe Met Arg Ala Leu Cys Val Val Cys Val Tyr Leu Arg Arg
20 25 30
Arg Ile Lys Tyr Glu Arg Phe Leu Gly Trp Glu Leu Arg Cys Lys Ile
35 40 45
Trp Gly *
50

<210> 1874
<211> 503
<212> PRT
<213> Homo sapiens

<400> 1874
Met Ser Leu Val Leu Leu Ser Leu Ala Ala Leu Cys Arg Ser Ala Val
1 5 10 15
Pro Arg Glu Pro Thr Val Gln Cys Gly Ser Glu Thr Gly Pro Ser Pro
20 25 30

Glu Trp Met Leu Gln His Asp Leu Ile Pro Gly Asp Leu Arg Asp Leu
35 40 45
Arg Val Glu Pro Val Thr Thr Ser Val Ala Thr Gly Asp Tyr Ser Ile
50 55 60
Leu Met Asn Val Ser Trp Val Leu Arg Ala Asp Ala Ser Ile Arg Leu
65 70 75 80
Leu Lys Ala Thr Lys Ile Cys Val Thr Gly Lys Ser Asn Phe Gln Ser
85 90 95
Tyr Ser Cys Val Arg Cys Asn Tyr Thr Glu Ala Phe Gln Thr Gln Thr
100 105 110
Arg Pro Ser Gly Gly Lys Trp Thr Phe Ser Tyr Ile Gly Phe Pro Val
115 120 125
Glu Leu Asn Thr Val Tyr Phe Ile Gly Ala His Asn Ile Pro Asn Ala
130 135 140
Asn Met Asn Glu Asp Gly Pro Ser Met Ser Val Asn Phe Thr Ser Pro
145 150 155 160
Gly Cys Leu Asp His Ile Met Lys Tyr Lys Lys Lys Cys Val Lys Ala
165 170 175
Gly Ser Leu Trp Asp Pro Asn Ile Thr Ala Cys Lys Lys Asn Glu Glu
180 185 190
Thr Val Glu Val Asn Phe Thr Thr Thr Pro Leu Gly Asn Arg Tyr Met
195 200 205
Ala Leu Ile Gln His Ser Thr Ile Ile Gly Phe Ser Gln Val Phe Glu
210 215 220
Pro His Gln Lys Lys Gln Thr Arg Ala Ser Val Val Ile Pro Val Thr
225 230 235 240
Gly Asp Ser Glu Gly Ala Thr Val Gln Leu Thr Pro Tyr Phe Pro Thr
245 250 255
Cys Gly Ser Asp Cys Ile Arg His Lys Gly Thr Val Val Leu Cys Pro
260 265 270
Gln Thr Gly Val Pro Phe Pro Leu Asp Asn Asn Lys Ser Lys Pro Gly
275 280 285
Gly Trp Leu Pro Leu Leu Leu Leu Ser Leu Leu Val Ala Thr Trp Val
290 295 300
Leu Val Ala Gly Ile Tyr Leu Met Trp Arg His Glu Arg Ile Lys Lys
305 310 315 320
Thr Ser Phe Ser Thr Thr Thr Leu Leu Pro Pro Ile Lys Val Leu Val
325 330 335
Val Tyr Pro Ser Glu Ile Cys Phe His His Thr Ile Cys Tyr Phe Thr
340 345 350
Glu Phe Leu Gln Asn His Cys Arg Ser Glu Val Ile Leu Glu Lys Trp
355 360 365
Gln Lys Lys Lys Ile Ala Glu Met Gly Pro Val Gln Trp Leu Ala Thr
370 375 380
Gln Lys Lys Ala Ala Asp Lys Val Val Phe Leu Leu Ser Asn Asp Val
385 390 395 400
Asn Ser Val Cys Asp Gly Thr Cys Gly Lys Ser Glu Gly Ser Pro Ser
405 410 415
Glu Asn Ser Gln Asp Leu Phe Pro Leu Ala Phe Asn Leu Phe Cys Ser
420 425 430
Asp Leu Arg Ser Gln Ile His Leu His Lys Tyr Val Val Val Tyr Phe
435 440 445
Arg Glu Ile Asp Thr Lys Asp Asp Tyr Asn Ala Leu Ser Val Cys Pro
450 455 460
Lys Tyr His Leu Met Lys Asp Ala Thr Ala Phe Cys Ala Glu Leu Leu
465 470 475 480
His Val Lys Gln Gln Val Ser Ala Gly Lys Arg Ser Gln Ala Cys His
485 490 495
Asp Gly Cys Cys Ser Leu *

500 502

<210> 1875
<211> 158
<212> PRT
<213> Homo sapiens

<221> misc_feature
<222> (1)...(158)
<223> Xaa = any amino acid or nothing

<400> 1875
Met Xaa Pro Pro Thr Arg Pro Arg Thr Arg Gly Val Gly Ile Phe Tyr
1 5 10 15
Phe Val Ile Tyr Ile Ile Ile Ser Phe Leu Val Val Val Asn Met Tyr
20 25 30
Ile Ala Val Ile Leu Glu Asn Phe Ser Val Ala Thr Glu Glu Ser Thr
35 40 45
Glu Pro Leu Ser Glu Asp Asp Phe Glu Met Phe Tyr Glu Val Trp Glu
50 55 60
Lys Phe Asp Pro Asp Ala Thr Gln Phe Ile Glu Phe Ser Lys Leu Ser
65 70 75 80
Asp Phe Ala Ala Ala Leu Asp Pro Pro Leu Leu Ile Ala Lys Pro Asn
85 90 95
Lys Val Gln Leu Ile Ala Met Asp Leu Pro Met Val Ser Gly Asp Arg
100 105 110
Ile His Cys Leu Asp Ile Leu Phe Ala Phe Thr Lys Arg Val Leu Gly
115 120 125
Glu Ser Gly Glu Met Asp Ser Leu Arg Ser Gln Met Glu Glu Arg Phe
130 135 140
Met Ser Ala Asn Pro Ser Lys Val Ser Tyr Glu Pro Ile Thr
145 150 155 158

<210> 1876
<211> 106
<212> PRT
<213> Homo sapiens

<400> 1876
Met Gly Asn Arg Ala Val Ile Ile Ala Arg Gln Leu Ser Ser Val His
1 5 10 15
Thr Leu Ile Cys Asn Phe Phe Trp Leu Leu Leu Arg Thr Thr Gly Gly
20 25 30
Asp Leu Asp Ser Leu Lys Cys Ser Tyr Glu Ser Ile Gly Leu Asn Ser
35 40 45
Ile Ser Thr His Glu Phe Ile Cys Thr Trp Gln Arg Arg Leu Asn Phe
50 55 60
Ser Phe Val Met Ser Phe Lys Pro Leu Phe Arg Ala Ser Pro His Ser
65 70 75 80
Tyr Leu Leu Ile Ile Gly Ser Gln Leu His Glu Thr Phe Asn Leu Gly
85 90 95
Ser Ile Ser Ser Glu Glu Lys Cys Ser *
100 105

<210> 1877
<211> 241
<212> PRT
<213> Homo sapiens

<221> misc_feature
<222> (1)...(241)
<223> Xaa = any amino acid or nothing

<400> 1877
Met Leu Trp Ala Leu Trp Pro Arg Trp Leu Ala Asp Lys Met Leu Pro
1 5 10 15
Leu Leu Gly Ala Val Leu Leu Gln Lys Arg Glu Lys Arg Gly Pro Leu
20 25 30
Trp Arg His Trp Arg Arg Glu Thr Tyr Pro Tyr Tyr Asp Leu Gln Val
35 40 45
Lys Val Leu Arg Ala Thr Asn Ile Arg Gly Thr Asp Leu Leu Ser Lys
50 55 60
Ala Asp Cys Tyr Val Gln Leu Trp Leu Pro Thr Ala Ser Pro Ser Pro
65 70 75 80
Ala Gln Thr Arg Ile Val Ala Asn Cys Ser Asp Pro Glu Trp Asn Glu
85 90 95
Thr Phe His Tyr Gln Ile His Gly Ala Val Lys Asn Val Leu Glu Leu
100 105 110
Thr Leu Tyr Asp Lys Asp Ile Leu Gly Ser Asp Gln Leu Ser Leu Leu
115 120 125
Leu Phe Asp Leu Arg Ser Leu Lys Cys Gly Gln Pro His Lys His Thr
130 135 140
Phe Pro Leu Asn His Gln Asp Ser Gln Glu Leu Gln Val Glu Phe Val
145 150 155 160
Leu Glu Lys Ser Gln Glu Pro Ala Ser Glu Val Ile Thr Asn Gly Val
165 170 175
Leu Gly Ala His Pro Trp Leu Arg Met Lys Gly Met Ile Leu Gly Glu
180 185 190
Gly Arg Ala Pro Arg Gln Gln His Gly Gln Ser Trp Glu Gly Gly Val
195 200 205
Gly Pro Ser Pro Leu Ser Xaa Xaa Xaa Asn Thr Gly Gly Lys Ile Val
210 215 220
Gly Phe Trp Glu Glu Met Ala Asn Gly Thr Gly Ala Pro Pro Arg Pro
225 230 235 240
Pro
241

<210> 1878
<211> 50
<212> PRT
<213> Homo sapiens

<400> 1878
Met Leu Leu Met Leu Leu Phe Arg Cys Cys Ser Ser Lys Asp Leu Trp
1 5 10 15
Pro Val Leu Ile Ala His Leu Val Pro Gln Gly Gly Gln Glu Gly Asn

20 25 30
Val Gly Glu Gln Thr Lys Gly Lys Ser Asn Arg Val Leu Pro Val Phe
35 40 45
Leu *
49

<210> 1879
<211> 56
<212> PRT
<213> Homo sapiens

<400> 1879
Met Cys Ser Ala Phe Ser Ser Phe Trp Trp Val Pro Pro Leu Ala Gly
1 5 10 15
Ser Gly Val Lys Leu Gln Thr Phe Thr Ala Ser Val Thr Ala His Lys
20 25 30
Arg Ser Thr Asp Pro Lys Ser Glu Gln Gln Leu Asp Leu Ser Gln Arg
35 40 45
Thr Lys Glu Gln Ser Leu Thr Lys
50 55 56

<210> 1880
<211> 161
<212> PRT
<213> Homo sapiens

<221> misc_feature
<222> (1)...(161)
<223> Xaa = any amino acid or nothing

<400> 1880
Met Pro Ser Ala Ser Leu Leu Val Asn Leu Leu Ser Ala Leu Leu Ile
1 5 10 15
Leu Phe Val Phe Gly Glu Thr Glu Ile Arg Phe Thr Gly Gln Thr Glu
20 25 30
Phe Val Val Asn Glu Thr Ser Thr Thr Val Ile Arg Leu Ile Ile Glu
35 40 45
Arg Ile Gly Glu Pro Ala Asn Val Thr Ala Ile Val Ser Leu Tyr Gly
50 55 60
Glu Asp Ala Gly Asp Phe Phe Asp Thr Tyr Ala Ala Ala Phe Ile Pro
65 70 75 80
Ala Gly Glu Thr Asn Arg Thr Val Tyr Ile Ala Val Cys Asp Asp Asp
85 90 95
Leu Pro Glu Pro Asp Glu Thr Phe Ile Phe His Leu Thr Leu Gln Lys
100 105 110
Pro Ser Ala Asn Val Lys Leu Gly Trp Pro Arg Thr Val Thr Val Thr
115 120 125
Ile Leu Ser Asn Gly Gln Met Ala Phe Trp Glu Phe Ile Phe Ile Leu
130 135 140
Asn Ile Gly Leu Pro Pro Pro Ile Pro Pro Ser Gly Xaa Leu Lys Ala
145 150 155 160
Pro
161

<210> 1881
<211> 130
<212> PRT
<213> Homo sapiens

<400> 1881
```
Met Gly Ile Tyr Gln Met Tyr Leu Cys Phe Leu Leu Ala Val Leu Leu
1               5                   10                  15
Gln Leu Tyr Val Ala Thr Glu Ala Ile Leu Ile Ala Leu Val Gly Ala
            20                  25                  30
Thr Pro Ser Tyr His Trp Asp Leu Ala Glu Leu Leu Pro Asn Gln Ser
        35                  40                  45
His Gly Asn Gln Ser Ala Gly Glu Asp Gln Ala Phe Gly Asp Trp Leu
    50                  55                  60
Leu Thr Ala Asn Gly Ser Glu Ile His Lys His Val His Phe Ser Ser
65                  70                  75                  80
Ser Phe Thr Ser Ile Ala Ser Glu Trp Phe Leu Ile Ala Asn Arg Ser
                85                  90                  95
Tyr Lys Val Ser Ala Ala Ser Ser Phe Phe Phe Ser Gly Val Phe Val
            100                 105                 110
Gly Val Ile Ser Phe Gly Gln Leu Ser Asp Arg Phe Gly Arg Lys Lys
        115                 120                 125
Val Tyr
    130
```

<210> 1882
<211> 108
<212> PRT
<213> Homo sapiens

<400> 1882
```
Met Leu Trp Phe Ser Gly Val Gly Ala Leu Ala Glu Arg Tyr Cys Arg
1               5                   10                  15
Arg Ser Pro Gly Ile Thr Cys Cys Val Leu Leu Leu Leu Asn Cys Ser
            20                  25                  30
Gly Val Pro Met Ser Leu Ala Ser Ser Phe Leu Thr Gly Ser Val Ala
        35                  40                  45
Lys Cys Glu Asn Glu Gly Glu Val Leu Gln Ile Pro Phe Ile Thr Asp
    50                  55                  60
Asn Pro Cys Ile Met Cys Val Cys Leu Asn Lys Glu Val Thr Cys Lys
65                  70                  75                  80
Arg Glu Lys Cys Pro Val Leu Ser Arg Asp Cys Ala Leu Ala Ile Lys
                85                  90                  95
Gln Arg Gly Ala Cys Cys Glu Gln Cys Lys Gly Cys
            100                 105         108
```

<210> 1883
<211> 88
<212> PRT
<213> Homo sapiens

<400> 1883
Met Leu Phe Tyr Leu Val Ser Val Cys Leu Cys Val Ala Val Ile Val
1 5 10 15
Ala Phe Gln Leu Thr Ala Phe Thr Phe Arg Lys Asn Leu Ala Ala Thr
20 25 30
Ala Leu Leu Leu Ser Leu Phe Gly Tyr Ala Thr Leu Pro Trp Met Tyr
35 40 45
Leu Met Ser Arg Ile Phe Ser Ser Ser Asp Val Ala Phe Ile Ser Tyr
50 55 60
Val Ser Leu Asn Phe Ile Phe Gly Leu Cys Thr Met Leu Ile Thr Ile
65 70 75 80
Met Pro Arg Leu Leu Ala Ile Ile
85 88

<210> 1884
<211> 116
<212> PRT
<213> Homo sapiens

<400> 1884
Met Cys Trp Ala Arg Cys Trp Thr Arg Trp Asn Thr Cys Thr Ile Trp
1 5 10 15
Thr Ser Ser Thr Asp Pro Phe Arg Lys Cys Trp Met Ala Pro Glu Ala
20 25 30
Leu Asn Phe Ser Phe Ser His Lys Ser Asp Ile Trp Ser Leu Gly Cys
35 40 45
Ile Ile Leu Asp Met Thr Ser Cys Ser Phe Met Asp Gly Thr Glu Ala
50 55 60
Met His Leu Arg Lys Ser Leu Arg Gln Ser Pro Gly Ser Leu Lys Ala
65 70 75 80
Val Leu Lys Thr Met Glu Glu Lys Gln Ile Pro Asp Val Glu Thr Phe
85 90 95
Arg Asn Leu Leu Pro Leu Met Leu Gln Ile Asp Pro Ser Asp Arg Ile
100 105 110
Thr Ile Lys *
115

<210> 1885
<211> 115
<212> PRT
<213> Homo sapiens

<400> 1885
Met Ser Glu Arg Val Glu Arg Asn Trp Ser Thr Gly Gly Trp Leu Leu
1 5 10 15
Ala Leu Cys Leu Ala Trp Leu Trp Thr His Leu Thr Leu Ala Ala Leu
20 25 30
Gln Pro Pro Thr Ala Thr Val Leu Val Gln Gln Gly Thr Cys Glu Val
35 40 45
Ile Ala Ala His Arg Cys Cys Asn Arg Asn Arg Ile Glu Glu Arg Ser
50 55 60

Gln Thr Val Lys Cys Ser Cys Phe Ser Gly Gln Val Ala Gly Thr Thr
65 70 75 80
Arg Ala Lys Pro Ser Cys Val Asp Asp Leu Leu Leu Ala Ala His Cys
85 90 95
Ala Arg Arg Asp Pro Arg Ala Ala Leu Arg Leu Leu Leu Pro Gln Pro
100 105 110
Pro Ser Ser
115

<210> 1886
<211> 357
<212> PRT
<213> Homo sapiens

<400> 1886
Met Ile Leu Ser Leu Leu Phe Ser Leu Gly Gly Pro Leu Gly Trp Gly
1 5 10 15
Leu Leu Gly Ala Trp Ala Gln Ala Ser Ser Thr Ser Leu Ser Asp Leu
20 25 30
Gln Ser Ser Arg Thr Pro Gly Val Trp Lys Ala Glu Ala Glu Asp Thr
35 40 45
Gly Lys Asp Pro Val Gly Arg Asn Trp Cys Pro Tyr Pro Met Ser Lys
50 55 60
Leu Val Thr Leu Leu Ala Leu Cys Lys Thr Glu Lys Phe Leu Ile His
65 70 75 80
Ser Gln Gln Pro Cys Pro Gln Gly Ala Pro Asp Cys Gln Lys Val Lys
85 90 95
Val Met Tyr Arg Met Ala His Lys Pro Val Tyr Gln Val Lys Gln Lys
100 105 110
Val Leu Thr Ser Leu Ala Trp Arg Cys Cys Pro Gly Tyr Thr Gly Pro
115 120 125
Asn Cys Glu His His Asp Ser Met Ala Ile Pro Glu Pro Ala Asp Pro
130 135 140
Gly Asp Ser His Gln Glu Pro Gln Asp Gly Pro Val Ser Phe Lys Pro
145 150 155 160
Gly His Leu Ala Ala Val Ile Asn Glu Val Glu Val Gln Gln Glu Gln
165 170 175
Gln Glu His Leu Leu Gly Asp Leu Gln Asn Asp Val His Arg Val Ala
180 185 190
Asp Ser Leu Pro Gly Leu Trp Lys Ala Leu Pro Gly Asn Leu Thr Ala
195 200 205
Ala Val Met Glu Ala Asn Gln Thr Gly His Glu Phe Pro Asp Arg Ser
210 215 220
Leu Glu Gln Val Leu Leu Pro His Val Asp Thr Phe Leu Gln Val His
225 230 235 240
Phe Ser Pro Ile Trp Arg Ser Phe Asn Gln Ser Leu His Ser Leu Thr
245 250 255
Gln Ala Ile Arg Asn Leu Ser Leu Asp Val Glu Ala Asn Arg Gln Ala
260 265 270
Ile Ser Arg Val Gln Asp Ser Ala Val Ala Arg Ala Asp Phe Gln Glu
275 280 285
Leu Gly Ala Lys Phe Glu Ala Lys Val Gln Glu Asn Thr Gln Arg Val
290 295 300
Gly Gln Leu Arg Gln Asp Val Glu Asp Arg Leu His Ala Gln His Phe
305 310 315 320
Thr Leu His Arg Ser Ile Ser Glu Leu Gln Ala Asp Val Asp Thr Lys

                325                 330                 335
Leu Lys Arg Leu His Lys Ala Gln Glu Ala Pro Gly Thr Asn Gly Ser
            340                 345                 350
Leu Val Leu Glu Arg
        355     357

<210> 1887
<211> 86
<212> PRT
<213> Homo sapiens

<400> 1887
Met Leu Cys Ser Arg Leu Gly Thr Thr Ala Ser Trp Arg Arg Leu Gly
 1               5                  10                  15
Ile Arg Ala Trp Ala Pro Leu Leu Leu Leu Phe Pro Trp Asp Trp His
            20                  25                  30
Phe Ile Leu Ser Phe Ser Ser Arg Pro Trp Ala Gly Thr Leu Leu Ala
        35                  40                  45
Pro His Asp Val Ile Met Gly Ser Ser Thr Phe Pro Gln Ser Cys Gln
    50                  55                  60
Ala Glu Ala Gly Pro Arg His Ala Trp Pro Thr Gly Arg Phe Ser Arg
65                  70                  75                  80
Arg Leu Arg Arg Val  *
                85

<210> 1888
<211> 48
<212> PRT
<213> Homo sapiens

<400> 1888
Met Ser Val Arg Arg Ala Leu Thr Pro Ser Ala Leu Gly Leu Val Phe
 1               5                  10                  15
Ile Leu Gln Ile Phe Ala His Gly Leu Pro Gly Pro Gly Pro Cys His
            20                  25                  30
Leu Gly Pro Gly Ile Cys Leu Arg Ile Cys Gln Cys Ala Leu Asn  *
        35                  40                  45      47

<210> 1889
<211> 79
<212> PRT
<213> Homo sapiens

<400> 1889
Met Ser Val Val Met Leu Ser Tyr Leu Leu Ser Ala Phe Phe Ser Gln
 1               5                  10                  15
Ala Asn Thr Ala Ala Leu Cys Thr Ser Leu Val Tyr Met Ile Ser Phe
            20                  25                  30
Leu Pro Tyr Ile Val Leu Leu Val Leu His Asn Gln Leu Ser Phe Val
        35                  40                  45

```
Asn Gln Thr Phe Leu Cys Leu Leu Ser Thr Thr Ala Phe Gly Gln Gly
    50                  55                  60
Val Phe Phe Ile Thr Phe Leu Glu Gly Gln Glu Thr Gly Ile His
 65                 70                  75              79
```

<210> 1890
<211> 251
<212> PRT
<213> Homo sapiens

<400> 1890

```
Met Asn Val Ile Tyr Phe Pro Leu His Leu Phe Val Val Tyr Ser Arg
  1               5                  10                  15
Ala Tyr Thr Ser Leu Val Leu Val Gly Cys Thr Asn Leu Cys Ala Val
             20                  25                  30
Leu Phe Ala Arg Cys Leu Asp Asp His Leu Val Ser Leu Arg Met Ser
         35                  40                  45
Gly Ser Arg Lys Glu Phe Asp Val Lys Gln Ile Leu Lys Ile Arg Trp
    50                  55                  60
Arg Trp Phe Gly His Gln Ala Ser Ser Pro Asn Ser Thr Val Asp Ser
 65                 70                  75                  80
Gln Gln Gly Glu Phe Trp Asn Arg Gly Gln Thr Gly Ala Asn Gly Gly
                 85                  90                  95
Arg Lys Phe Leu Asp Pro Cys Ser Leu Gln Leu Pro Leu Ala Ser Ile
            100                 105                 110
Gly Tyr Arg Arg Ser Ser Gln Leu Asp Phe Gln Asn Ser Pro Ser Trp
        115                 120                 125
Pro Met Ala Ser Thr Ser Glu Val Pro Ala Phe Glu Phe Thr Ala Glu
    130                 135                 140
Asp Cys Gly Gly Ala His Trp Leu Asp Arg Pro Glu Val Asp Asp Gly
145                 150                 155                 160
Thr Ser Glu Glu Glu Asn Glu Ser Asp Ser Ser Ser Cys Arg Thr Ser
                165                 170                 175
Asn Ser Ser Gln Thr Leu Ser Ser Cys His Thr Met Glu Pro Cys Thr
            180                 185                 190
Ser Asp Glu Phe Phe Gln Ala Leu Asn His Ala Glu Gln Thr Phe Lys
        195                 200                 205
Lys Met Glu Asn Tyr Leu Arg His Lys Gln Leu Cys Asp Val Ile Leu
    210                 215                 220
Val Ala Gly Asp Arg Arg Ile Pro Ala His Arg Leu Val Leu Ser Ser
225                 230                 235                 240
Val Ser Asp Tyr Phe Ala Gly Met Phe Thr Asn
                245                 250 251
```

<210> 1891
<211> 117
<212> PRT
<213> Homo sapiens

<221> misc_feature
<222> (1)...(117)
<223> Xaa = any amino acid or nothing

<400> 1891
Met Leu Ile Asp Val Phe Phe Phe Leu Phe Leu Phe Ala Xaa Trp Met
1 5 10 15
Val Ala Phe Gly Val Ala Arg Gln Gly Ile Leu Arg Gln Asn Glu Gln
20 25 30
Arg Trp Arg Trp Ile Phe Arg Ser Val Ile Tyr Glu Pro Tyr Leu Ala
35 40 45
Met Phe Gly Gln Val Pro Ser Asp Val Asp Gly Thr Thr Tyr Asp Phe
50 55 60
Ala His Cys Thr Phe Thr Gly Asn Glu Ser Lys Pro Leu Cys Val Glu
65 70 75 80
Leu Asp Glu His Asn Leu Pro Arg Phe Pro Glu Trp Ile Thr Ile Pro
85 90 95
Leu Val Cys Ile Tyr Met Leu Ser Thr Asn Ile Leu Leu Val Asn Leu
100 105 110
Leu Val Ala Met Phe
115 117

<210> 1892
<211> 103
<212> PRT
<213> Homo sapiens

<400> 1892
Met Leu Cys His Pro His Val His His His Leu Val Cys Leu Leu Ala
1 5 10 15
Thr Leu Thr Phe Ser Leu Asn Ala Ser Cys Ala Glu Gln Thr Phe His
20 25 30
Ser Gln Gln Ser Asn Gly Glu Phe Met Ala Thr Leu Pro Ser Ile Ser
35 40 45
Lys Gln Phe Gly Val Ile Val Trp Lys Pro Gln Arg Lys Asp Val Ile
50 55 60
Arg Leu Pro Val Ala Leu Ser Phe Ser Ser Gly Ala Arg Leu Ala Phe
65 70 75 80
Thr Cys Leu Arg Lys Ile Ser Gly Phe Arg Ala Leu Ile Trp Gly Glu
85 90 95
Asp Lys Gly Trp Asp Leu *
100 102

<210> 1893
<211> 77
<212> PRT
<213> Homo sapiens

<221> misc_feature
<222> (1)...(77)
<223> Xaa = any amino acid or nothing

<400> 1893
Met Leu Ala Ala Gly Val Thr Ser Ala Ala Gly Leu Ala Leu Ala Phe
1 5 10 15
Ser Gly Asp Tyr Leu Lys Ala Phe Ile Asp Val Pro Thr Val Pro Ala
20 25 30

```
Ala Leu Val Phe Leu Leu Leu Val Gly Leu Leu Asn Ala Arg Gly Ile
        35                  40                  45
Lys Glu Ser Met Arg Ala Xaa Val Val Met Thr Val Val Glu Val Thr
    50                  55                  60
Gly Leu Val Leu Val Val Val Leu Ala Leu Val Pro Gly
65                  70                  75      77



     <210> 1894
     <211> 46
     <212> PRT
     <213> Homo sapiens


     <400> 1894
Met Trp Ala Ala Ser Trp Cys Leu Ser Leu Trp Cys Cys Trp Val Trp
 1               5                  10                  15
Ser Gly Thr Ser Glu Ser Ile Thr Ala Asn Ser Ser Gln His Leu Pro
            20                  25                  30
Leu Ser Pro Trp Trp Glu Ser Pro Ser Ser Ser Ala Ser  *
        35                  40                  45



     <210> 1895
     <211> 162
     <212> PRT
     <213> Homo sapiens


     <400> 1895
Met Thr Ala Trp Arg Arg Phe Gln Ser Leu Leu Leu Leu Leu Gly Leu
 1               5                  10                  15
Leu Val Leu Cys Ala Arg Leu Leu Thr Ala Ala Lys Gly Gln Asn Cys
            20                  25                  30
Gly Gly Leu Val Gln Gly Pro Asn Gly Thr Ile Glu Ser Pro Gly Phe
        35                  40                  45
Pro His Gly Tyr Pro Asn Tyr Ala Asn Cys Thr Trp Ile Ile Ile Thr
    50                  55                  60
Gly Glu Arg Asn Arg Ile Gln Leu Ser Phe His Thr Phe Ala Leu Glu
65                  70                  75                  80
Glu Asp Phe Asp Ile Leu Ser Val Tyr Asp Gly Gln Pro Gln Gln Gly
                85                  90                  95
Asn Leu Lys Val Arg Leu Ser Gly Phe Gln Leu Pro Ser Ser Ile Val
            100                 105                 110
Ser Thr Gly Ser Ile Leu Thr Leu Trp Phe Thr Thr Asp Phe Ala Val
        115                 120                 125
Ser Ala Gln Gly Phe Lys Ala Leu Tyr Glu Gly Arg Arg Leu Val Val
    130                 135                 140
Phe Cys Thr Cys Ile His Cys Pro Asn Asp Leu Ile His Ala Thr Leu
145                 150                 155                 160
Asp  *
161



     <210> 1896
     <211> 60
```

<212> PRT
<213> Homo sapiens

<400> 1896
Met Leu Ser Leu Pro Cys Gly Trp Leu Cys Thr Ala Ile Gly Leu Pro
1 5 10 15
Thr Met Phe Gly Tyr Ile Ile Cys Gly Val Leu Leu Gly Pro Ser Gly
20 25 30
Leu Asn Ser Ile Lys Val Arg Thr Lys Leu Asp Cys Phe Gly Ile Cys
35 40 45
Leu Thr Glu Tyr Lys Lys Arg Ile His Glu Asp *
50 55 59

<210> 1897
<211> 49
<212> PRT
<213> Homo sapiens

<400> 1897
Met Leu Ile Val Gln Phe Ile Phe Glu Leu Val Ser Ser Ile Leu Val
1 5 10 15
Ser Asn Val Lys Asp Met Leu Asp Phe Glu Ser Gly Phe Cys Ser Lys
20 25 30
Ile Leu Ser Tyr Phe Phe Ser Ser Pro Arg Tyr Arg Leu Pro Phe Leu
35 40 45 48
*

<210> 1898
<211> 52
<212> PRT
<213> Homo sapiens

<400> 1898
Met Thr Trp Ala Gly Leu Phe Leu Phe Leu Arg Val Gly Ser Pro Asn
1 5 10 15
Arg Lys Trp Ala Ala Ser Gly Gly Ser Gly Gly Asp Gly Val Asp Gly
20 25 30
Glu Asp Trp Ser Leu Ala Arg Ser His Pro Gln Ser Pro Leu Leu Leu
35 40 45
Leu Leu Leu *
50 51

<210> 1899
<211> 112
<212> PRT
<213> Homo sapiens

<400> 1899

Met Ala Ile Pro Ser Val Val Ile Ser Gly Leu Ala Val Leu Leu Val
1 5 10 15
Ala Met Ala Leu Pro Ser Leu Ser Gly Ser Glu Ala Ile Lys Ser Met
20 25 30
Thr Ile Pro Gly Leu Val Val Pro Thr Val Val Arg Phe Met Ala Val
35 40 45
Pro Gly Leu Ile Val Pro Ala Val Ala Lys Phe Thr Val Leu Pro Asp
50 55 60
Leu Thr Val Pro Thr Glu Asp Lys Ser Leu Ala Val Pro Ser Leu Ile
65 70 75 80
Ser Arg Ala Gly Asn Ser Val Pro Val Ser Ser Trp Asp Val Phe Gly
85 90 95
Val Ala Lys Leu Ile Ala Lys Leu Gly Leu Leu Ala Ala Ile Val Ala
100 105 110 112

<210> 1900
<211> 128
<212> PRT
<213> Homo sapiens

<400> 1900
Met Arg Val Tyr Gly Thr Cys Thr Leu Val Leu Met Ala Leu Val Val
1 5 10 15
Phe Val Gly Val Lys Tyr Val Asn Lys Leu Ala Leu Val Phe Leu Ala
20 25 30
Cys Val Val Leu Ser Ile Leu Ala Ile Tyr Ala Gly Val Ile Lys Ser
35 40 45
Ala Phe Asp Pro Pro Asp Ile Pro Val Cys Leu Leu Gly Asn Arg Thr
50 55 60
Leu Ser Arg Arg Ser Phe Asp Ala Cys Val Lys Ala Tyr Gly Ile His
65 70 75 80
Asn Asn Ser Ala Thr Ser Ala Leu Trp Gly Leu Phe Cys Asn Gly Ser
85 90 95
Gln Pro Ser Ala Ala Cys Asp Glu Tyr Phe Ile Gln Asn Asn Val Thr
100 105 110
Glu Ile Gln Gly Ile Pro Gly Ala Ala Ser Gly Val Phe Leu Glu Asn
115 120 125 128

<210> 1901
<211> 68
<212> PRT
<213> Homo sapiens

<400> 1901
Met Glu Leu Leu Lys Leu Leu Leu Thr Cys Phe Ser Glu Ala Met Tyr
1 5 10 15
Leu Pro Pro Ala Pro Glu Ser Gly Ser Thr Asn Pro Trp Val Gln Phe
20 25 30
Phe Cys Ser Thr Glu Asn Arg His Ala Leu Pro Leu Phe Thr Ser Leu

```
        35                  40                  45
Leu Asn Thr Val Cys Ala Tyr Asp Pro Val Glu Tyr Gly Ile Pro Tyr
    50                  55                  60
Asn His Leu Tyr
 65         68
```

<210> 1902
<211> 127
<212> PRT
<213> Homo sapiens

<400> 1902
```
Met Tyr Phe Ser Ser Leu Phe Pro Tyr Val Val Leu Ala Cys Phe Leu
 1               5                  10                  15
Val Arg Gly Leu Leu Leu Arg Gly Ala Val Asp Gly Ile Leu His Met
            20                  25                  30
Phe Thr Pro Lys Leu Asp Lys Met Leu Asp Pro Gln Val Trp Arg Glu
        35                  40                  45
Ala Ala Thr Gln Val Phe Ser Ala Leu Gly Leu Gly Phe Gly Gly Val
    50                  55                  60
Ile Ala Phe Ser Ser Tyr Asn Lys Gln Asp Asn Asn Cys His Phe Asp
 65                 70                  75                  80
Ala Ala Leu Val Ser Phe Ile Asn Phe Phe Thr Ser Val Leu Ala Thr
                85                  90                  95
Leu Val Val Phe Ala Val Leu Gly Phe Lys Ala Asn Ile Met Asn Glu
            100                 105                 110
Lys Cys Val Val Glu Asn Ala Glu Lys Ile Leu Gly Tyr Arg Val
        115                 120                 125     127
```

<210> 1903
<211> 83
<212> PRT
<213> Homo sapiens

<400> 1903
```
Met Trp Lys Phe Val Ser Pro Leu Cys Met Ala Val Leu Thr Thr Ala
 1               5                  10                  15
Ser Ile Ile Gln Leu Gly Val Thr Pro Pro Gly Tyr Ser Ala Trp Ile
            20                  25                  30
Lys Glu Glu Ala Ala Glu Arg Tyr Leu Tyr Phe Pro Asn Trp Ala Met
        35                  40                  45
Ala Pro Leu Ile Thr Leu Ile Val Val Ala Thr Leu Pro Ile Pro Val
    50                  55                  60
Val Phe Val Leu Arg His Phe His Leu Ile Cys Asp Gly Ser Asn Thr
 65                 70                  75                  80
Pro Cys Ile
        83
```

<210> 1904
<211> 129
<212> PRT

<213> Homo sapiens

<400> 1904
Met Lys Met Phe Val Ala His Gly Phe Tyr Ala Ala Lys Phe Val Val
1               5                   10                  15
Ala Ile Gly Ser Val Ala Gly Leu Thr Val Ser Leu Leu Gly Ser Leu
            20                  25                  30
Phe Pro Met Pro Arg Val Ile Tyr Ala Met Ala Gly Asp Gly Leu Leu
        35                  40                  45
Phe Arg Phe Leu Ala His Val Ser Ser Tyr Thr Glu Thr Pro Val Val
    50                  55                  60
Ala Cys Ile Val Ser Gly Phe Leu Ala Ala Leu Leu Ala Leu Leu Val
65                  70                  75                  80
Ser Leu Arg Asp Leu Ile Glu Met Met Ser Ile Gly Thr Leu Leu Ala
                85                  90                  95
Tyr Thr Leu Val Ser Val Cys Val Leu Leu Leu Arg His His Pro Glu
            100                 105                 110
Ser Asp Ile Asp Gly Phe Val Lys Phe Leu Ser Glu Glu His Thr Cys
        115                 120                 125
Ser
129

<210> 1905
<211> 93
<212> PRT
<213> Homo sapiens

<400> 1905
Met Gly Leu Leu Met Met Ile Leu Gly Gln Ile Phe Leu Asn Gly Asn
1               5                   10                  15
Gln Ala Lys Glu Ala Glu Ile Trp Glu Met Leu Trp Arg Met Gly Val
            20                  25                  30
Gln Arg Glu Arg Arg Leu Ser Ile Phe Gly Asn Pro Lys Arg Leu Leu
        35                  40                  45
Ser Val Glu Phe Val Trp Gln Arg Tyr Leu Asp Tyr Arg Pro Val Thr
    50                  55                  60
Asp Cys Lys Pro Val Glu Tyr Glu Phe Phe Trp Gly Pro Arg Ser His
65                  70                  75                  80
Leu Glu Thr Thr Lys Met Lys Ile Leu Lys Phe Met Ala
                85                  90          93

<210> 1906
<211> 66
<212> PRT
<213> Homo sapiens

<400> 1906
Met Thr Ile Gly Phe Leu Phe Pro Met Leu Ser Ile Ala Tyr Leu Ile
1               5                   10                  15
Ser Pro Arg Ser Asn Leu Gly Leu Phe Ile Lys Lys Pro Phe Ile Lys
            20                  25                  30
Phe Ile Cys His Thr Ala Ser Tyr Leu Thr Phe Leu Ser Met Leu Leu

```
        35                  40                  45
Leu Ala Ser Gln His Ile Val Arg Thr Asp Leu His Val Gln Gly Pro
    50                  55                  60
Cys Ile
65  66
```

<210> 1907
<211> 105
<212> PRT
<213> Homo sapiens

<400> 1907

```
Met Leu Gln Leu Gly Pro Phe Leu Tyr Trp Thr Phe Leu Ala Ala Phe
 1               5                  10                  15
Glu Gly Thr Val Phe Phe Phe Gly Thr Tyr Phe Leu Phe Gln Thr Ala
            20                  25                  30
Ser Leu Glu Glu Asn Gly Lys Val Tyr Gly Asn Trp Thr Phe Gly Thr
        35                  40                  45
Ile Val Phe Thr Val Leu Val Phe Thr Val Thr Leu Lys Leu Ala Leu
    50                  55                  60
Asp Thr Arg Phe Trp Thr Trp Ile Asn His Phe Val Ile Trp Gly Ser
65                  70                  75                  80
Leu Ala Phe Tyr Val Phe Phe Ser Phe Phe Trp Gly Gly Ile Ile Trp
                85                  90                  95
Pro Phe Leu Lys Gln Gln Arg Met Ala
            100                 105
```

<210> 1908
<211> 46
<212> PRT
<213> Homo sapiens

<400> 1908

```
Met Gly Phe Leu Val Leu Lys Gln Pro Met Leu Val Ala Lys Val Phe
 1               5                  10                  15
Pro Thr Leu Ala Gly Val Glu Ile Ile Leu Phe Thr Leu Lys Gly Phe
            20                  25                  30
Pro Ile Leu Gly Ile Pro Val Gln Leu Pro Pro Thr Val  *
        35                  40                  45
```

<210> 1909
<211> 139
<212> PRT
<213> Homo sapiens

<400> 1909

```
Met Ile Gln Ala Leu Gly Gly Phe Phe Thr Tyr Phe Val Ile Leu Ala
 1               5                  10                  15
Glu Asn Gly Phe Leu Pro Ile His Leu Leu Gly Leu Arg Glu Asp Trp
            20                  25                  30
```

```
Asp Asp Arg Trp Ile Asn Asp Val Glu Asp Ser Tyr Gly Gln Gln Trp
        35                  40                  45
Thr Tyr Glu Gln Arg Lys Ile Val Glu Phe Thr Cys His Thr Ala Phe
    50                  55                  60
Phe Val Ser Ile Val Gly Val Gln Trp Ala Asp Leu Val Ile Cys Lys
65                  70                  75                  80
Thr Arg Arg Asn Ser Val Phe Gln Pro Gly Met Lys Asn Lys Ile Leu
                85                  90                  95
Ile Phe Gly Leu Phe Glu Glu Thr Ala Leu Ala Ala Phe Leu Ser Tyr
            100                 105                 110
Cys Pro Gly Met Gly Val Ala Leu Lys Met Tyr Pro Leu Lys Pro Thr
        115                 120                 125
Trp Arg Val Cys Ala Phe Pro Tyr Ser Leu Leu
    130                 135             139
```

<210> 1910
<211> 104
<212> PRT
<213> Homo sapiens

<400> 1910

```
Met Glu Gly Trp Phe Ala Val Leu Ser Thr Ala Asn Asp Val Leu Gly
1               5                   10                  15
Ala Pro Trp Asn Trp Leu Tyr Phe Ile Pro Leu Leu Ile Ile Gly Ala
            20                  25                  30
Phe Phe Val Pro Thr Leu Val Leu Gly Val Leu Ser Gly Asp Phe Ala
        35                  40                  45
Lys Glu Arg Glu Arg Val Glu Thr Arg Arg Ala Phe Met Lys Leu Arg
    50                  55                  60
Arg Gln Gln Gln Ile Glu Arg Glu Leu Asn Gly Tyr Arg Val Trp Ile
65                  70                  75                  80
Ala Lys Ala Glu Glu Val Met Leu Ala Glu Glu Asn Leu Tyr Pro Ser
                85                  90                  95
His Ala Arg Pro Val Asn Pro  *
            100         103
```

<210> 1911
<211> 116
<212> PRT
<213> Homo sapiens

<400> 1911

```
Met Ala Val Ala Val Leu Leu Cys Gly Cys Ile Val Ala Thr Val Ser
1               5                   10                  15
Phe Phe Trp Glu Glu Ser Leu Thr Gln His Val Ala Gly Leu Leu Phe
            20                  25                  30
Leu Met Thr Gly Ile Phe Cys Thr Ile Ser Leu Cys Thr Tyr Ala Ala
        35                  40                  45
Ser Ile Ser Tyr Asp Leu Asn Arg Leu Pro Lys Leu Ile Tyr Ser Leu
    50                  55                  60
Pro Ala Asp Val Glu His Gly Tyr Ser Trp Ser Ile Phe Cys Ala Trp
65                  70                  75                  80
Cys Ser Leu Gly Phe Ile Val Ala Ala Gly Gly Leu Cys Ile Ala Tyr
```

```
                85                      90                      95
Pro Phe Ile Ser Arg Thr Lys Ile Ala Gln Leu Lys Ser Gly Arg Asp
            100                     105                     110
Ser Thr Val  *
        115
```

<210> 1912
<211> 105
<212> PRT
<213> Homo sapiens

<400> 1912

```
Met Gln Leu Lys Thr Pro Ser Gly Gln Val Leu Ser Phe Cys Ile Leu
 1               5                  10                  15
Gln Leu Phe Pro Phe Thr Ser Glu Ser Lys Arg Met Gly Val Ile Val
            20                  25                  30
Arg Asp Glu Ser Thr Ala Glu Ile Thr Phe Tyr Met Lys Gly Ala Asp
        35                  40                  45
Val Ala Met Ser Pro Ile Val Gln Tyr Asn Asp Trp Leu Glu Glu Glu
    50                  55                  60
Cys Gly Asn Met Ala Arg Glu Gly Leu Arg Thr Leu Val Val Ala Lys
65                  70                  75                  80
Lys Ala Leu Thr Glu Glu Gln Tyr Gln Asp Phe Glu Ser Arg Tyr Thr
                85                  90                  95
Gln Ala Lys Leu Ser Met His Thr Lys
            100                 105
```

<210> 1913
<211> 141
<212> PRT
<213> Homo sapiens

<400> 1913

```
Met Leu Val Tyr Val Trp Ser Arg Arg Ser Pro Arg Val Arg Val Asn
 1               5                  10                  15
Phe Phe Gly Leu Leu Thr Phe Gln Ala Pro Phe Leu Pro Trp Ala Leu
            20                  25                  30
Met Gly Phe Ser Leu Leu Leu Gly Asn Ser Ile Leu Val Asp Leu Leu
        35                  40                  45
Gly Ile Ala Val Gly His Ile Tyr Tyr Phe Leu Glu Asp Val Phe Pro
    50                  55                  60
Asn Gln Pro Gly Arg Gln Glu Ala Pro Ala Asp Pro Trp Ala Phe Leu
65                  70                  75                  80
Lys Leu Leu Leu Gly Cys Pro Cys Arg Arg Pro Gln Leu Thr Cys Pro
                85                  90                  95
Ser Leu Arg Asn Ser Gln Asp Pro Ile Cys His Pro Arg Ser Ser Asp
            100                 105                 110
Pro His Pro Gly Ala Arg Pro Lys Arg Leu Leu Ala Ala Ser Ile Leu
        115                 120                 125
Pro Met Thr Pro Thr Trp Gly Arg Lys Asn Pro Ser  *
    130                 135                 140
```

<210> 1914
<211> 556
<212> PRT
<213> Homo sapiens

<400> 1914
Met Lys Lys Val Leu Leu Leu Leu Trp Lys Thr Val Leu Cys Thr Leu
1 5 10 15
Gly Gly Phe Glu Glu Leu Gln Ser Met Lys Ala Glu Lys Arg Ser Ile
20 25 30
Leu Gly Leu Pro Pro Leu Pro Glu Asp Ser Ile Lys Val Ile Arg Asn
35 40 45
Met Arg Ala Ala Ser Pro Pro Ala Ser Ala Ser Asp Leu Ile Glu Gln
50 55 60
Gln Gln Lys Arg Gly Arg Arg Glu His Lys Ala Leu Ile Lys Gln Asp
65 70 75 80
Asn Leu Asp Ala Phe Asn Glu Arg Asp Pro Tyr Lys Ala Asp Asp Ser
85 90 95
Arg Glu Glu Glu Glu Glu Asn Asp Asp Asp Asn Ser Leu Glu Gly Glu
100 105 110
Thr Phe Pro Leu Glu Arg Asp Glu Val Met Pro Pro Pro Leu Gln His
115 120 125
Pro Gln Thr Asp Arg Leu Thr Cys Pro Lys Gly Leu Pro Trp Ala Pro
130 135 140
Lys Val Arg Glu Lys Asp Ile Glu Met Phe Leu Glu Ser Ser Arg Ser
145 150 155 160
Lys Phe Ile Gly Tyr Thr Leu Gly Ser Asp Thr Asn Thr Val Val Gly
165 170 175
Leu Pro Arg Pro Ile His Glu Ser Ile Lys Thr Leu Lys Gln His Lys
180 185 190
Tyr Thr Ser Ile Ala Glu Val Gln Ala Gln Met Glu Glu Glu Tyr Leu
195 200 205
Arg Ser Pro Leu Ser Gly Gly Glu Glu Glu Val Glu Gln Val Pro Ala
210 215 220
Glu Thr Leu Tyr Gln Gly Leu Leu Pro Ser Leu Pro Gln Tyr Met Ile
225 230 235 240
Ala Leu Leu Lys Ile Leu Leu Ala Ala Ala Pro Thr Ser Lys Ala Lys
245 250 255
Thr Asp Ser Ile Asn Ile Leu Ala Asp Val Leu Pro Glu Glu Met Pro
260 265 270
Thr Thr Val Leu Gln Ser Met Lys Leu Gly Val Asp Val Asn Arg His
275 280 285
Lys Glu Val Ile Val Lys Ala Ile Ser Ala Val Leu Leu Leu Leu Leu
290 295 300
Lys His Phe Lys Leu Asn His Val Tyr Gln Phe Glu Tyr Met Ala Gln
305 310 315 320
His Leu Val Phe Ala Asn Cys Ile Pro Leu Ile Leu Lys Phe Phe Asn
325 330 335
Gln Asn Ile Met Ser Tyr Ile Thr Ala Lys Asn Ser Ile Ser Val Leu
340 345 350
Asp Tyr Pro His Cys Val Val His Glu Leu Pro Glu Leu Thr Ala Glu
355 360 365
Ser Leu Glu Ala Gly Asp Ser Asn Gln Phe Cys Trp Arg Asn Leu Phe
370 375 380
Ser Cys Ile Asn Leu Leu Arg Ile Leu Asn Lys Leu Thr Lys Trp Lys
385 390 395 400
His Ser Arg Thr Met Met Leu Val Val Phe Lys Ser Ala Pro Ile Leu

405 410 415
Lys Arg Ala Leu Lys Val Lys Gln Ala Met Met Gln Leu Tyr Val Leu
420 425 430
Lys Leu Leu Lys Val Gln Thr Lys Tyr Leu Gly Arg Gln Trp Arg Lys
435 440 445
Ser Asn Met Lys Thr Met Ser Ala Ile Tyr Gln Lys Val Arg His Arg
450 455 460
Leu Asn Asp Asp Trp Ala Tyr Gly Asn Asp Leu Asp Ala Arg Pro Trp
465 470 475 480
Asp Phe Gln Ala Glu Glu Cys Ala Leu Arg Ala Asn Ile Glu Arg Phe
485 490 495
Asn Ala Arg Arg Tyr Asp Arg Ala His Ser Asn Pro Asp Phe Leu Pro
500 505 510
Val Asp Asn Cys Leu Gln Ser Val Leu Gly Gln Arg Val Asp Leu Pro
515 520 525
Glu Asp Phe Gln Met Asn Tyr Asp Leu Trp Leu Glu Arg Glu Val Phe
530 535 540
Ser Lys Pro Ile Ser Trp Glu Glu Leu Leu Gln *
545 550 555

<210> 1915
<211> 212
<212> PRT
<213> Homo sapiens

<400> 1915
Met Phe Leu Val Ala Val Trp Trp Arg Phe Gly Ile Leu Ser Ile Cys
1 5 10 15
Met Leu Cys Val Gly Leu Val Leu Gly Phe Leu Ile Ser Ser Val Thr
20 25 30
Phe Phe Thr Pro Leu Gly Asn Leu Lys Ile Phe His Asp Asp Gly Val
35 40 45
Phe Trp Val Thr Phe Ser Cys Ile Ala Ile Leu Ile Pro Val Val Phe
50 55 60
Met Gly Cys Leu Arg Ile Leu Asn Ile Leu Thr Cys Gly Val Ile Gly
65 70 75 80
Ser Tyr Ser Val Val Leu Ala Ile Asp Ser Tyr Trp Ser Thr Ser Leu
85 90 95
Ser Tyr Ile Thr Leu Asn Val Leu Lys Arg Ala Leu Asn Lys Asp Phe
100 105 110
His Arg Ala Phe Thr Asn Val Pro Phe Gln Thr Asn Asp Phe Ile Ile
115 120 125
Leu Ala Val Trp Gly Met Leu Ala Val Ser Gly Ile Thr Leu Gln Ile
130 135 140
Arg Arg Glu Arg Gly Arg Pro Phe Phe Pro Pro His Pro Tyr Lys Leu
145 150 155 160
Trp Lys Gln Glu Arg Glu Arg Arg Val Thr Asn Ile Leu Asp Pro Ser
165 170 175
Tyr His Ile Pro Pro Leu Arg Glu Arg Leu Tyr Gly Arg Leu Thr Gln
180 185 190
Ile Lys Gly Leu Phe Gln Lys Glu Gln Pro Ala Gly Glu Arg Thr Pro
195 200 205
Leu Leu Leu *
210 211

<210> 1916
<211> 172
<212> PRT
<213> Homo sapiens

<400> 1916
Met Cys Thr Pro Val Arg Val Ser Ile Val Cys Val Met Gly Ala Val
1 5 10 15
Gly Ala Val Trp Thr Ala Pro Leu Pro Leu Pro Trp Ala Pro Thr Pro
20 25 30
Ser Ile His Leu Arg Glu Glu Gly Ala Ala Phe Pro Phe Cys Gly Val
35 40 45
Cys Val Leu Arg Pro Arg Arg Ser Lys Trp Arg Ser Trp Asp Val Asn
50 55 60
Leu Gly Pro Arg Arg Arg Gly Leu Leu Gly Cys Gly Pro Cys Pro Ser
65 70 75 80
Gly Lys Pro Arg Val His Leu Gln Arg Thr Arg Ser Gly Ala Gly Ala
85 90 95
Glu Ala Gly Gly Leu Pro Thr Arg Gly Ser Met Arg Gly Cys Pro Phe
100 105 110
Leu Gly Ser Ser Ala Ala Lys Cys Ser Leu Leu Leu Arg Pro Pro Ser
115 120 125
Arg Gly Glu Ala Ser Pro Trp Leu Pro Glu Phe Met Thr His Pro Val
130 135 140
His His Gln Gln Leu Ala Cys Gly Ser Gly Trp Leu Gly Thr Lys His
145 150 155 160
Pro Gly Gly Thr Cys Ala Leu Gly Ser Thr Met *
165 170 171

<210> 1917
<211> 72
<212> PRT
<213> Homo sapiens

<400> 1917
Met Leu Arg Trp Gly Phe Leu Glu Ile Leu Phe Leu Arg Ser Trp Phe
1 5 10 15
His Ser Trp Ile Cys Leu Leu Pro Thr Pro Gln Leu Pro Pro Asn Gly
20 25 30
Ala Ser Ala Gly Ser Gln Asp Glu Gly Ser Arg Arg Arg Leu Ser Leu
35 40 45
Glu Val Arg Gly Leu Met Asn His Val Pro Asn Leu Cys Val Ala Phe
50 55 60
Leu Ser Ile Val Ser Ile Ser *
65 70 71

<210> 1918
<211> 88
<212> PRT
<213> Homo sapiens

<400> 1918
Met Thr Ser Leu Met Phe Leu Trp Arg Ala Leu Leu Glu Thr Ile Ser
1 5 10 15
Thr Asn Met Thr Phe Ser Leu Pro Leu Ala Ala Val Val Arg Ala Trp
20 25 30
Met Lys Pro Thr Gly Ser Gly Met Phe Leu Tyr Gln Tyr Leu Pro Val
35 40 45
Val Lys Ser Ser Gln Ala Val Phe Pro Val Val Ile Glu Ile Ser Ser
50 55 60
Ile Ser Gly Ser Ile Leu Pro Lys Phe Pro Met Leu Ser Leu Met Ser
65 70 75 80
Leu His Thr Gly Ser Ile Ile *
85 87

<210> 1919
<211> 54
<212> PRT
<213> Homo sapiens

<400> 1919
Met Leu Gly Pro Phe Ser Ser Leu Phe Leu Leu Leu Trp Ser Phe Thr
1 5 10 15
Arg Phe Cys Ile His Phe Tyr Leu Ala Pro Ser His His Cys Leu Thr
20 25 30
Ala Ala Leu Leu Pro Phe Ser Leu His Pro Leu Tyr Ser Ser Leu Ser
35 40 45
Leu Ser Arg Ser Gln *
50 53

<210> 1920
<211> 114
<212> PRT
<213> Homo sapiens

<400> 1920
Met His Pro Pro Leu Thr Pro Pro Thr Pro Leu Cys Leu Trp Leu Arg
1 5 10 15
Leu Leu Lys Ala Gln Ile Leu Ser Tyr Pro Val Pro Arg Phe Glu Thr
20 25 30
His Ser Leu Ile Ser Arg Cys Ser Gln Val Pro Pro Thr Phe Leu Trp
35 40 45
Asp Ile Lys Lys Gly Val Arg Gly Gln Arg Glu Pro Ser Gly Pro Leu
50 55 60
Leu Pro Tyr Thr Leu His Cys Pro Phe Ser Pro His Gln Asn Ala Gln
65 70 75 80
Arg Arg Cys Asp Asp Ala Thr Glu Asp Tyr Ala Thr Trp Ser Asn Arg
85 90 95
Ser Gly Gln His Asp Gln Leu Ser Arg Gly Cys Leu Leu Pro Phe Leu
100 105 110
Leu *
113

<210> 1921
<211> 139
<212> PRT
<213> Homo sapiens

<400> 1921
```
Met Val Tyr Leu Tyr Ile Tyr Leu Asp Leu Phe Gln Phe Leu Ile Thr
 1               5                  10                  15
Val Leu Gln Gly Phe Leu Phe Val Phe Glu Met Glu Phe His Ser Cys
            20                  25                  30
Arg Pro Gly Gln Ser Ala Met Met Gln Ser Gln Leu Ala Ala Thr Ser
        35                  40                  45
Ala Ser Arg Val Gln Val Ile Leu Val Val Ser Ala Pro Gln Glu Ala
    50                  55                  60
Gly Thr Thr Gly Ala Arg His His Val Gln Leu Ile Phe Val Phe Leu
 65                  70                  75                  80
Leu Glu Met Gly Phe Cys His Val Gly Gln Ala Gly Leu Glu Leu Leu
                85                  90                  95
Asn Ser Gly Asp Pro Pro Thr Ser Ala Ser Gln Ser Ala Gly Ile Arg
            100                 105                 110
Gly Val Asn His Cys Ala Pro Pro Ile Asn Ser Leu Leu Thr Phe Gln
        115                 120                 125
Ser Phe Ile His Leu Glu Cys Ile Val Ile  *
    130                 135         138
```

<210> 1922
<211> 52
<212> PRT
<213> Homo sapiens

<400> 1922
```
Met Trp Leu Ser Phe Pro Lys Leu Phe Ile Pro Leu Ser Ile Phe Leu
 1               5                  10                  15
Val Phe Leu Leu Met Ala Asn Ser Phe Arg Ile Phe Lys Ser Lys Asn
            20                  25                  30
Ile Phe Ile Ser Leu Leu Phe Trp Asn Asp Thr Phe Ala Gly Cys Ile
        35                  40                  45
Phe Leu Thr  *
    50  51
```

<210> 1923
<211> 71
<212> PRT
<213> Homo sapiens

<400> 1923
```
Met Val Ser His Cys Ile Phe Cys Asn Leu Leu Phe Ser Leu Leu Thr
 1               5                  10                  15
Val Phe Leu Arg Leu Leu His Val Asp Thr Cys His Leu Phe Ile Arg
            20                  25                  30
Phe Asn Cys Cys Lys Ile Phe Phe Cys Gln Asp Ile Leu Gln Leu Ile
```

```
        35                  40                  45
Tyr Leu Leu Phe Phe Leu Trp Thr Phe Lys Leu Phe Ser Gly Phe Thr
    50                  55                  60
Leu Lys Ile Ile Gln Gln  *
 65                  70
```

<210> 1924
<211> 187
<212> PRT
<213> Homo sapiens

<400> 1924

```
Met Leu Phe Ile Gln Tyr Leu Leu Pro Cys Leu Leu Leu Ser Ala Glu
  1               5                  10                  15
Leu Ser Gly Thr Phe Phe Leu Tyr Asn Thr Cys His Leu His Val Pro
             20                  25                  30
Cys Cys His Ser Leu Val Pro Thr Gly Pro Pro Ser Leu Ser Ser His
        35                  40                  45
Phe Gln Ser Arg Gly Leu Cys Ala Pro Cys Ala Ser Ile Ala Asp Ser
    50                  55                  60
Gly Ile Ala Asp Ser Gly Gly Asn Asn Leu Asn Phe Val Gly Ala Gly
 65                  70                  75                  80
Gly Val Ala Ser Gly His Leu Leu Ser Pro Leu Leu Gly Pro Gln Ser
                 85                  90                  95
Ser Pro Cys Pro His Cys Pro Arg Gly Gly Arg Leu Pro Ser Gln Pro
            100                 105                 110
Leu Pro Leu Cys Ser Ala Arg Ser Trp Ala Gln Glu Ala Leu Arg Leu
        115                 120                 125
Pro Ser Ser Ala Gln Leu Cys Pro Cys His Pro Leu Pro Arg Gly Leu
    130                 135                 140
Gly Pro Val Ser Pro Ser Gly Leu Leu Ala Asn Ile Ser Tyr Arg His
145                 150                 155                 160
Asn Trp Leu Leu Gly Ser Trp Pro Gly Trp Leu Ile Trp Gly Gly Lys
                165                 170                 175
Asn Arg Gly Gly Leu Asn Ser Phe Leu Ala  *
            180                 185 186
```

<210> 1925
<211> 50
<212> PRT
<213> Homo sapiens

<400> 1925

```
Met Leu Ser Phe Leu Val Val Phe Gln Leu Val Leu Leu Arg Phe Ser
  1               5                  10                  15
Gly Arg His Ser His His Gln Leu Ile Thr Ile Thr Phe Pro Leu Phe
             20                  25                  30
Gln Trp Leu Tyr Phe Phe Phe Phe Met Phe Phe Cys Thr Gly Trp Lys
        35                  40                  45
Phe  *
 49
```

<210> 1926
<211> 47
<212> PRT
<213> Homo sapiens

<400> 1926
Met Gly Arg Tyr Arg Cys Ala Ser Leu Leu Phe Cys Phe Leu Leu Leu
1 5 10 15
Phe Phe Phe Phe Trp Leu Trp Val Arg Asp Ile Phe Lys Leu Ala Gln
20 25 30
Lys Gly Arg Gly Trp Ser Leu Asp Pro His Val Ser Ile Thr *
35 40 45 46

<210> 1927
<211> 149
<212> PRT
<213> Homo sapiens

<400> 1927
Met Ala Thr Gly Leu Leu Ala Phe Leu Gly Leu Ala Ala Gly Gly Gln
1 5 10 15
Thr Leu Cys Pro Ala Gly Glu Leu Pro Gly His Ala Arg Ala Gln Ala
20 25 30
Ser Gly Ala Pro Gly Ser Val Leu Ile Ala Val Pro Gly Arg Arg Arg
35 40 45
Val His Thr Cys Gly Pro Gly Pro Ala Ala Pro Ser Thr Arg Gly Glu
50 55 60
Cys Pro Pro Pro Ala Leu Gly His Thr Arg Pro Ala Arg Pro Arg Pro
65 70 75 80
Val Leu Leu Arg Pro Ser Cys Ser Pro Gly Ala Arg Gly Ala Gly Thr
85 90 95
Trp Cys Cys Ala Pro Ala Thr Gly His Ser Ala Pro Arg Gly Cys Pro
100 105 110
Pro Ala Arg Ala Ala Pro Thr Gly Ser Ala Thr Pro Ala Pro Pro Pro
115 120 125
Ala Ala Cys Ala Ala Phe His Ser Ala Trp Ser Val Pro Pro Ala Gly
130 135 140
Arg Gln Gln Gly *
145 148

<210> 1928
<211> 446
<212> PRT
<213> Homo sapiens

<400> 1928
Met Ser Leu Trp Asn Gln Leu Val Val Pro Val Leu Phe Met Val Phe
1 5 10 15
Trp Leu Val Leu Phe Ala Leu Gln Ile Tyr Ser Tyr Phe Ser Thr Arg
20 25 30
Asp Gln Pro Ala Ser Arg Glu Arg Leu Leu Phe Leu Phe Leu Thr Ser

35 40 45
Ile Ala Glu Cys Cys Ser Thr Pro Tyr Ser Leu Leu Gly Leu Val Phe
50 55 60
Thr Val Ser Phe Val Ala Leu Gly Val Leu Thr Leu Cys Lys Phe Tyr
65 70 75 80
Leu Gln Gly Tyr Arg Ala Phe Met Asn Asp Pro Ala Met Asn Arg Gly
85 90 95
Met Thr Glu Gly Val Thr Leu Leu Ile Leu Ala Val Gln Thr Gly Leu
100 105 110
Ile Glu Leu Gln Val Val His Arg Ala Phe Leu Leu Ser Ile Ile Leu
115 120 125
Phe Ile Val Val Ala Ser Ile Leu Gln Ser Met Leu Glu Ile Ala Asp
130 135 140
Pro Ile Val Leu Ala Leu Gly Ala Ser Arg Asp Lys Ser Leu Trp Lys
145 150 155 160
His Phe Arg Ala Val Ser Leu Cys Leu Phe Leu Leu Val Phe Pro Ala
165 170 175
Tyr Met Ala Tyr Met Ile Cys Gln Phe Phe His Met Asp Phe Trp Leu
180 185 190
Leu Ile Ile Ile Ser Ser Ser Ile Leu Thr Ser Leu Gln Val Leu Gly
195 200 205
Thr Leu Phe Ile Tyr Val Leu Phe Met Val Glu Glu Phe Arg Lys Glu
210 215 220
Pro Val Glu Asn Met Asp Asp Val Ile Tyr Tyr Val Asn Gly Thr Tyr
225 230 235 240
Arg Leu Leu Glu Phe Leu Val Ala Leu Cys Val Val Ala Tyr Gly Val
245 250 255
Ser Glu Thr Ile Phe Gly Glu Trp Thr Val Met Gly Ser Met Ile Ile
260 265 270
Phe Ile His Ser Tyr Tyr Asn Val Trp Leu Arg Ala Gln Leu Gly Trp
275 280 285
Lys Ser Phe Leu Leu Arg Arg Asp Ala Val Asn Lys Ile Lys Ser Leu
290 295 300
Pro Ile Ala Thr Lys Glu Gln Leu Glu Lys His Asn Asp Ile Cys Ala
305 310 315 320
Ile Cys Tyr Gln Asp Met Lys Ser Ala Val Ile Thr Pro Cys Ser His
325 330 335
Phe Phe His Ala Gly Cys Leu Lys Lys Trp Leu Tyr Val Gln Glu Thr
340 345 350
Cys Pro Leu Cys His Cys His Leu Lys Asn Ser Ser Gln Leu Pro Gly
355 360 365
Leu Gly Thr Glu Pro Val Leu Gln Pro His Ala Gly Ala Glu Gln Asn
370 375 380
Val Met Phe Gln Glu Gly Thr Glu Pro Pro Gly Gln Glu His Thr Pro
385 390 395 400
Gly Thr Arg Ile Gln Glu Gly Ser Arg Asp Asn Asn Glu Tyr Ile Ala
405 410 415
Arg Arg Pro Asp Asn Gln Glu Gly Ala Phe Asp Pro Lys Glu Tyr Pro
420 425 430
His Ser Ala Lys Asp Glu Ala His Pro Val Glu Ser Ala *
435 440 445

<210> 1929
<211> 120
<212> PRT
<213> Homo sapiens

<400> 1929
Met Val Leu Pro Leu Pro Trp Leu Ser Arg Tyr His Phe Leu Arg Leu
1 5 10 15
Leu Leu Pro Ser Trp Ser Leu Ala Pro Gln Gly Ser His Gly Cys Cys
20 25 30
Ser Gln Asn Pro Lys Ala Ser Met Glu Glu Gln Thr Asn Ser Arg Gly
35 40 45
Asn Gly Lys Met Thr Ser Pro Pro Arg Gly Pro Gly Thr His Arg Thr
50 55 60
Ala Glu Leu Ala Arg Ala Glu Glu Leu Leu Glu Gln Gln Leu Glu Leu
65 70 75 80
Tyr Gln Ala Leu Leu Glu Gly Gln Glu Gly Ala Trp Glu Ala Gln Ala
85 90 95
Leu Val Leu Lys Ile His Lys Leu Lys Glu Gln Met Arg Arg His Gln
100 105 110
Glu Ser Leu Gly Gly Gly Ala *
115 119

<210> 1930
<211> 122
<212> PRT
<213> Homo sapiens

<400> 1930
Met Thr Trp Leu Val Leu Leu Gly Thr Leu Leu Cys Met Leu Arg Val
1 5 10 15
Gly Leu Gly Thr Pro Asp Ser Glu Gly Phe Pro Pro Arg Ala Leu His
20 25 30
Asn Cys Pro Tyr Lys Cys Ile Cys Ala Ala Asp Leu Leu Ser Cys Thr
35 40 45
Gly Leu Gly Leu Gln Asp Val Pro Ala Glu Leu Pro Ala Gly Thr Ala
50 55 60
Asp Leu Asp Leu Ser His Asn Ala Leu Gln Arg Met Arg Pro Gly Trp
65 70 75 80
Leu Ala Pro Leu Phe Gln Leu Arg Ala Leu His Leu Asp His Asn Glu
85 90 95
Leu His Ala Leu Asp Arg Gly Val Phe Val Asn Ala Ser Gly Leu Arg
100 105 110
Leu Leu Asp Leu Ser Ser Asn Ala Glu Phe
115 120 122

<210> 1931
<211> 73
<212> PRT
<213> Homo sapiens

<400> 1931
Met Ala Arg Ala Pro Ser Val Ala Leu Ala Gln Leu Trp Leu Ile Cys
1 5 10 15
Leu Cys Pro Glu Ser Leu Ala Ser Phe Val Gln Ala Val Pro Trp Lys
20 25 30
Val Leu Gln Pro Ser Ser Asn Arg Ser Thr Asp Cys Ser Pro His Met

35 40 45
Arg Pro Thr Cys Glu Thr Leu Gly Ser Arg Lys Ala Gln Asp Leu Gly
50 55 60
Ala Gly Tyr Tyr Val Ser Val His *
65 70 72

<210> 1932
<211> 68
<212> PRT
<213> Homo sapiens

<400> 1932
Met Lys Thr Val Phe Thr Lys Lys Leu Thr Ala Ala Leu Leu Ile Thr
1 5 10 15
Val Pro Asn Cys Lys Gln Pro Arg Cys Pro Ser Met Gly Glu Trp Leu
20 25 30
Asn Lys Leu Gln Tyr Ile His Thr Met Lys Tyr Tyr Ser Thr Ile Lys
35 40 45
Val Asn Tyr Trp Pro Gly Thr Val Ala His Thr Cys Asn Pro Ser Thr
50 55 60
Leu Gly Gly *
65 67

<210> 1933
<211> 47
<212> PRT
<213> Homo sapiens

<400> 1933
Met Gln Gln Arg Lys Met Arg Leu Val Trp Arg Ser Tyr Trp Ser Met
1 5 10 15
Val Gln Thr Pro Met Leu Trp Met Ala Thr Glu Ile Pro His Phe Thr
20 25 30
Gly Gln Pro Leu Arg Thr Met Leu Ser Val Cys Gly Leu Ser *
35 40 45 46

<210> 1934
<211> 86
<212> PRT
<213> Homo sapiens

<400> 1934
Met Cys Trp Ser Pro Leu Thr Gly Trp Ala Leu Ser Ser Ser Arg Cys
1 5 10 15
Arg Leu Ser Trp Pro Leu Thr Ser Phe Gly Ser Thr Ala Ser Cys Arg
20 25 30
Pro Thr Thr Gly Trp Arg Gly Leu Met Trp Leu Gln Ala Leu Ser Ser
35 40 45
Ser Gly Tyr Pro Ser Leu Cys Thr Leu Tyr Ser Glu Leu Leu Val Gln
50 55 60

Ala Val His Arg Lys Ala Gly Asp Thr Glu Val Gln Gln Ser Leu Leu
65 70 75 80
Leu Leu Leu Lys Lys *
85

<210> 1935
<211> 76
<212> PRT
<213> Homo sapiens

<400> 1935
Met Gly Glu Val Pro Lys Ala His Arg Leu Lys Leu Arg Trp Leu Phe
1 5 10 15
Pro Val Ser Leu Cys Arg Ala Pro Leu Leu Ser Thr Ala His Leu Ala
20 25 30
Leu Leu Leu Pro Cys Cys Leu Leu Cys Ser Ser Cys Tyr Tyr Phe Pro
35 40 45
Phe Leu Ser Leu Leu Pro Pro Trp Pro Asn Leu Phe His Arg Asn Ile
50 55 60
Thr Gly Pro Ala Arg His Ser Gly Ser Pro Leu *
65 70 75

<210> 1936
<211> 49
<212> PRT
<213> Homo sapiens

<400> 1936
Met Leu Leu Gln Thr Phe Val Thr Thr Cys Ile Ser Tyr Phe Tyr Trp
1 5 10 15
His Phe Asn Phe Val Trp Ile Gln Phe Asn Val Cys Arg Val Leu Ser
20 25 30
Phe Gln Pro Glu Arg Leu Thr Leu Ala Phe Leu Ile Gly Gln Val Tyr
35 40 45 48
*

<210> 1937
<211> 76
<212> PRT
<213> Homo sapiens

<400> 1937
Met Lys Gly Arg Phe Leu Phe Pro Leu Arg Leu Leu Leu Trp Met Cys
1 5 10 15
Leu His Leu Gln Arg Gln Ala Ser Glu Leu His Gln Pro Ser Met Pro
20 25 30
Gly Cys Pro Leu Thr Ser Ser Ser Arg Leu Phe Asp Asn Ala Gln Met
35 40 45
His Gln Phe Leu Asn Ile His Val Lys Phe Glu Asn Cys Thr Phe Gly

```
    50                  55                  60
Glu Ile Lys Phe Tyr Ile Gln Leu Ala Lys Lys Lys
 65                  70                  75  76
```

```
     <210> 1938
     <211> 191
     <212> PRT
     <213> Homo sapiens

     <400> 1938
Met Ala Asp Glu Lys Thr Phe Arg Ile Gly Phe Ile Val Leu Gly Leu
  1               5                  10                  15
Phe Leu Leu Ala Leu Gly Thr Phe Leu Met Ser His Asp Arg Pro Gln
             20                  25                  30
Val Tyr Gly Thr Phe Tyr Ala Met Gly Ser Val Met Val Ile Gly Gly
         35                  40                  45
Ile Ile Trp Ser Met Cys Gln Cys Tyr Pro Lys Ile Thr Phe Val Pro
     50                  55                  60
Ala Asp Ser Asp Phe Gln Gly Ile Leu Ser Pro Lys Ala Met Gly Leu
 65                  70                  75                  80
Leu Glu Asn Gly Leu Ala Ala Glu Met Lys Ser Pro Ser Pro Gln Pro
                 85                  90                  95
Pro Tyr Val Arg Leu Trp Glu Glu Ala Ala Tyr Asp Gln Ser Leu Pro
            100                 105                 110
Asp Phe Ser His Ile Gln Met Lys Val Met Ser Tyr Ser Glu Asp His
        115                 120                 125
Arg Ser Leu Leu Ala Pro Glu Met Gly Gln Pro Lys Leu Gly Thr Ser
    130                 135                 140
Asp Gly Gly Glu Gly Gly Pro Gly Asp Val Gln Ala Trp Met Glu Ala
145                 150                 155                 160
Ala Val Val Ile His Lys Gly Leu Asn Glu Ser Glu Gly Glu Arg Arg
                165                 170                 175
Leu Thr Gln Ser Trp Pro Gly Pro Leu Ala Cys Pro Gln Gly Pro
            180                 185                 190 191
```

```
     <210> 1939
     <211> 82
     <212> PRT
     <213> Homo sapiens

     <400> 1939
Met Val Arg Ser Ile Arg Leu Leu Phe Phe Phe Gly Trp Gly Phe Ser
  1               5                  10                  15
Thr Thr Gln Gln Pro Ser Leu Cys Gln Asn Ser Leu Met Phe Pro Asp
             20                  25                  30
Gly Ser Ser Phe Thr Pro Leu Ser Glu Ala Pro Lys Gly Ser Phe Pro
         35                  40                  45
Gly Val Trp Thr Thr His Ser Ser Leu Ser Pro Asp Thr Pro Pro Pro
     50                  55                  60
Trp Val His Ser Ala Gly Trp Val Gln Thr Lys Trp Asn Pro Trp Asn
 65                  70                  75                  80
Leu  *
 81
```

<210> 1940
<211> 101
<212> PRT
<213> Homo sapiens

<400> 1940
Met His Val Cys Leu His Ile Trp Gly Leu Gly Val Cys Val Phe Met
1 5 10 15
His Met Met Cys Ala Cys Val Gly Val Tyr Val Cys Pro Phe Met Arg
20 25 30
Tyr Gly Met Gln Ile Cys Ala Cys Ile His Ala His Ser Cys Ser Ala
35 40 45
Cys Val Cys Ser Cys Ile Trp Cys Met His Gly Cys Ser Tyr Leu Trp
50 55 60
Gly Thr Gly Ile Met His Val Cys Ser Ser Val Trp Gly Val Gly Ile
65 70 75 80
Pro Gly Leu Trp Pro Glu Ala Pro Leu Gln Asp Thr Ala Pro Cys Arg
85 90 95
Leu Pro Arg Gly *
100

<210> 1941
<211> 88
<212> PRT
<213> Homo sapiens

<400> 1941
Met Lys Ala Ser Val Leu Ser Pro Ser Phe Leu Leu Val Leu Trp Ser
1 5 10 15
Cys Phe Leu Ser Cys Ser Cys Met Glu Pro Gln Ser Gly Phe Pro Arg
20 25 30
Pro Ser Cys Phe Thr Val Gly Phe Leu Leu Arg Arg Arg Thr Lys Thr
35 40 45
Arg Arg Gln Lys Ala Thr Asn Thr Val Lys Met Arg Thr Thr Lys Ile
50 55 60
Leu Lys Ile Lys Ile Asp Lys Arg Arg Trp Pro Thr Arg Met Ser Ser
65 70 75 80
Lys Trp Asn Pro Lys Glu Trp *
85 87

<210> 1942
<211> 46
<212> PRT
<213> Homo sapiens

<400> 1942
Met Arg Ser Met Gly Phe Arg Ala Gln Gly Leu Pro Phe Gly Ile Arg
1 5 10 15
Gln Thr Trp Leu Arg Ile Leu Asp Leu Leu Leu Thr Cys Thr Leu Pro

```
            20                  25                  30
Phe Gly Ser Arg Asp Val Lys Trp Arg Cys Cys His Leu  *
        35                  40                  45
```

<210> 1943
<211> 155
<212> PRT
<213> Homo sapiens

<400> 1943

```
Met Phe Thr Leu Leu Val Leu Leu Ser Gln Leu Pro Thr Val Thr Leu
 1               5                  10                  15
Gly Phe Pro His Cys Ala Arg Gly Pro Lys Ala Ser Lys His Ala Gly
            20                  25                  30
Glu Glu Val Phe Thr Ser Lys Glu Glu Ala Asn Phe Phe Ile His Arg
        35                  40                  45
Arg Leu Leu Tyr Asn Arg Phe Asp Leu Glu Leu Phe Thr Pro Gly Asn
    50                  55                  60
Leu Glu Arg Glu Cys Asn Glu Glu Leu Cys Asn Tyr Glu Glu Ala Arg
65                  70                  75                  80
Glu Ile Phe Val Asp Glu Asp Lys Thr Ile Ala Phe Trp Gln Glu Tyr
                85                  90                  95
Ser Ala Lys Gly Pro Thr Thr Lys Ser Asp Gly Asn Arg Glu Lys Ile
            100                 105                 110
Asp Val Met Gly Leu Leu Thr Gly Leu Ile Ala Ala Gly Val Phe Leu
        115                 120                 125
Val Ile Phe Gly Leu Leu Gly Tyr Tyr Leu Cys Ile Thr Lys Cys Asn
    130                 135                 140
Arg Leu Gln His Pro Cys Ser Ser Ala Val Tyr
145                 150                 155
```

<210> 1944
<211> 61
<212> PRT
<213> Homo sapiens

<400> 1944

```
Met Cys Gln His Val Gln Leu Ile Phe Val Phe Phe Val Glu Thr Gly
 1               5                  10                  15
Phe His His Val Ala Gln Ala Gly Leu Lys Leu Leu Gly Ser Ser Asp
            20                  25                  30
Leu Pro Thr Ser Ala Ser Gln Ser Ala Gly Ile Lys Gly Ile Ser His
        35                  40                  45
His Val Gln Leu Lys Phe Leu Ile Ile Asn Asn Phe  *
    50                  55                  60
```

<210> 1945
<211> 79
<212> PRT
<213> Homo sapiens

<400> 1945
Met Gln Leu Ile Leu Trp Leu Pro Trp Tyr Val Asp Gln Thr Phe Cys
1 5 10 15
His Ser Val Leu Gln Cys Cys Cys Pro Gly Gln Leu Cys Gln Ser Phe
20 25 30
His Ser Asn Arg Asn Asp Ala Arg Leu Leu Gly Ala Lys Gln Ser Ile
35 40 45
Met Arg Arg Lys Arg Trp Leu Glu Pro Ser Val Arg Glu Cys Ala Pro
50 55 60
Gly Met Ile Leu Tyr Lys Ile Gln Ser Tyr Leu Lys Ile Gln *
65 70 75 78

<210> 1946
<211> 72
<212> PRT
<213> Homo sapiens

<400> 1946
Met Leu Arg Trp Gly Phe Leu Glu Ile Leu Phe Leu Arg Ser Trp Phe
1 5 10 15
His Ser Trp Ile Cys Leu Leu Pro Thr Pro Gln Leu Pro Pro Asn Gly
20 25 30
Ala Ser Ala Gly Ser Gln Asp Glu Gly Ser Arg Arg Arg Leu Ser Leu
35 40 45
Glu Val Arg Gly Leu Met Asn His Val Pro Asn Leu Cys Val Ala Phe
50 55 60
Leu Ser Ile Val Ser Ile Ser *
65 70 71

<210> 1947
<211> 56
<212> PRT
<213> Homo sapiens

<400> 1947
Met Trp Asn Val Ala Phe Leu Phe Gln Trp Phe Leu Ser Leu Lys Lys
1 5 10 15
Glu Gly Arg Ser Ser Val Glu Thr Lys Asp Arg Arg Ser Val Arg Asp
20 25 30
Leu Trp Gly Met Pro Lys Lys Met Val Ser Phe Gly Gly Glu Trp Leu
35 40 45
Arg Glu Gly Leu Arg Glu Val *
50 55

<210> 1948
<211> 48
<212> PRT
<213> Homo sapiens

<400> 1948
Met Ser Leu Leu Leu Pro Pro Leu Ala Leu Leu Leu Leu Leu Ala Ala
1 5 10 15
Leu Val Ala Pro Ala Thr Ala Ala Thr Ala Tyr Arg Pro Asp Trp Asn
20 25 30
Arg Leu Ser Gly Leu Thr Arg Ala Arg Val Glu Thr Cys Gly Gly *
35 40 45 47

<210> 1949
<211> 136
<212> PRT
<213> Homo sapiens

<400> 1949
Met Leu Leu Ala Thr Leu Leu Leu Leu Leu Leu Gly Gly Ala Leu Ala
1 5 10 15
His Pro Asp Arg Ile Ile Phe Pro Asn His Ala Cys Glu Asp Pro Pro
20 25 30
Ala Val Leu Leu Glu Val Gln Gly Thr Leu Gln Arg Pro Leu Val Arg
35 40 45
Asp Ser Arg Thr Ser Pro Ala Asn Cys Thr Trp Leu Ile Leu Gly Ser
50 55 60
Lys Glu Gln Thr Val Thr Ile Arg Phe Gln Lys Leu His Leu Ala Cys
65 70 75 80
Gly Ser Glu Arg Leu Thr Leu Arg Ser Pro Leu Gln Pro Leu Ile Ser
85 90 95
Leu Cys Glu Ala Pro Pro Ser Pro Leu Gln Leu Pro Gly Gly Asn Val
100 105 110
Thr Ile Thr Tyr Ser Tyr Ala Gly Ala Lys Arg Pro Gln Gly His Gly
115 120 125
Phe Phe Cys Phe Leu Lys Ala Lys
130 135 136

<210> 1950
<211> 78
<212> PRT
<213> Homo sapiens

<400> 1950
Met Trp Ile Tyr Phe Trp Thr Leu Asn Ser Val Pro Val Ile Tyr Met
1 5 10 15
Ser Thr Leu Met Ser Ile Pro His Tyr Phe Asp Tyr Cys Cys Phe Ile
20 25 30
Val Ser Asp Ile Met Leu Pro Glu Ile Thr Phe Ser Thr Phe Ile Leu
35 40 45
Leu Leu Met Val Ala Leu Ala Ile Arg Gly Pro Leu His Phe Arg Arg
50 55 60
His Phe Arg Ile Asn Leu Ser Ile Ala Thr Lys Asn Ala *
65 70 75 77

<210> 1951

<211> 89
<212> PRT
<213> Homo sapiens

<400> 1951
```
Met Val Cys Gly Ala Leu Met Trp Ile Met Leu Ile Leu Val Gly Leu
 1               5                  10                  15
Gly Phe Pro Phe Ile Met Glu Ala Leu Ser His Phe Leu Tyr Val Pro
            20                  25                  30
Phe Leu Gly Val Cys Val Cys Gly Ala Ile Tyr Thr Gly Leu Phe Leu
        35                  40                  45
Pro Glu Thr Lys Gly Lys Thr Phe Gln Glu Ile Ser Lys Glu Leu His
    50                  55                  60
Arg Leu Asn Phe Pro Arg Arg Ala Gln Gly Pro Thr Trp Arg Ser Leu
65                  70                  75                  80
Glu Val Ile Gln Ser Thr Glu Leu  *
                85          88
```

<210> 1952
<211> 47
<212> PRT
<213> Homo sapiens

<400> 1952
```
Met Thr Thr Ala Leu Ser Phe Met Val Ile Thr Val Leu Trp Val Leu
 1               5                  10                  15
Leu Leu His Leu Leu Ala Asn Ile Cys Ile Pro Arg Lys Cys Ser Phe
            20                  25                  30
Val Cys Phe Tyr Ile Asn Gly Ile Leu Leu His Ala Val Phe  *
        35                  40                  45  46
```

<210> 1953
<211> 56
<212> PRT
<213> Homo sapiens

<400> 1953
```
Met Lys Asn Leu Arg Leu Gly Glu Val Val Thr Leu Ser Trp Val Leu
 1               5                  10                  15
Val Val Glu Leu Glu Val Lys Ala Lys Ser Val Phe Leu Leu Ala Ile
            20                  25                  30
Leu Thr Thr Glu Phe Ser Leu Asn Gln Ser Leu Lys Met Phe Leu Gly
        35                  40                  45
Gln Glu Trp Trp Phe Thr Leu  *
    50                  55
```

<210> 1954
<211> 425
<212> PRT
<213> Homo sapiens

<400> 1954

Met Thr Leu Arg Pro Gly Thr Met Arg Leu Ala Cys Met Phe Ser Ser
1 5 10 15
Ile Leu Leu Phe Gly Ala Ala Gly Leu Leu Leu Phe Ile Ser Leu Gln
20 25 30
Asp Pro Thr Glu Leu Ala Pro Gln Gln Val Pro Gly Ile Lys Phe Asn
35 40 45
Ile Arg Pro Arg Gln Pro His His Asp Leu Pro Pro Gly Gly Ser Gln
50 55 60
Asp Gly Asp Leu Lys Glu Pro Thr Glu Arg Val Thr Arg Asp Leu Ser
65 70 75 80
Ser Gly Ala Pro Arg Gly Arg Asn Leu Pro Ala Pro Asp Gln Pro Gln
85 90 95
Pro Pro Leu Gln Arg Gly Thr Arg Leu Arg Leu Arg Gln Arg Arg Arg
100 105 110
Arg Leu Leu Ile Lys Lys Met Pro Ala Ala Ala Thr Ile Pro Ala Asn
115 120 125
Ser Ser Asp Ala Pro Phe Ile Arg Pro Gly Pro Gly Thr Leu Asp Gly
130 135 140
Arg Trp Val Ser Leu His Arg Ser Gln Gln Glu Arg Lys Arg Val Met
145 150 155 160
Gln Glu Ala Cys Ala Lys Tyr Arg Ala Ser Ser Ser Arg Arg Ala Val
165 170 175
Thr Pro Arg His Val Ser Arg Ile Phe Val Glu Asp Arg His Arg Val
180 185 190
Leu Tyr Cys Glu Val Pro Lys Ala Gly Cys Ser Asn Trp Lys Arg Val
195 200 205
Leu Met Val Leu Ala Gly Leu Ala Ser Ser Thr Ala Asp Ile Gln His
210 215 220
Asn Thr Val His Tyr Gly Ser Ala Leu Lys Arg Leu Asp Thr Phe Asp
225 230 235 240
Arg Gln Gly Ile Leu His Arg Leu Ser Thr Tyr Thr Lys Met Leu Phe
245 250 255
Val Arg Glu Pro Phe Glu Arg Leu Val Ser Ala Phe Arg Asp Lys Phe
260 265 270
Glu His Pro Asn Ser Tyr Tyr His Pro Val Phe Gly Lys Ala Ile Leu
275 280 285
Ala Arg Tyr Arg Ala Asn Ala Ser Arg Glu Ala Leu Arg Thr Gly Ser
290 295 300
Gly Val Arg Phe Pro Glu Phe Val Gln Tyr Leu Leu Asp Val His Arg
305 310 315 320
Pro Val Gly Met Asp Ile His Trp Asp His Val Ser Arg Leu Cys Ser
325 330 335
Pro Cys Leu Ile Asp Tyr Asp Phe Val Gly Lys Phe Glu Ser Met Glu
340 345 350
Asp Asp Ala Asn Phe Phe Leu Ser Leu Ile Arg Ala Pro Arg Asn Leu
355 360 365
Thr Phe Pro Arg Phe Lys Asp Arg His Ser Gln Glu Ala Arg Thr Thr
370 375 380
Ala Arg Ile Ala His Gln Tyr Phe Ala Gln Leu Ser Ala Leu Gln Arg
385 390 395 400
Gln Arg Thr Tyr Asp Phe Tyr Tyr Met Asp Tyr Leu Met Phe Asn Tyr
405 410 415
Ser Lys Pro Phe Ala Asp Leu Tyr *
420 424

<210> 1955
<211> 106
<212> PRT
<213> Homo sapiens

<400> 1955
Met Val Cys Phe Leu Phe Ile Thr Pro Leu Ala Ala Ile Ser Gly Trp
1 5 10 15
Leu Cys Leu Arg Gly Ala Gln Asp His Leu Arg Leu His Ser Gln Leu
20 25 30
Glu Ala Val Gly Leu Ile Ala Leu Thr Ile Ala Leu Phe Thr Ile Tyr
35 40 45
Val Leu Trp Thr Leu Val Ser Phe Arg Tyr His Cys Gln Leu Tyr Ser
50 55 60
Glu Trp Arg Lys Thr Asn Gln Lys Val Arg Leu Lys Ile Arg Glu Ala
65 70 75 80
Asp Ser Pro Glu Gly Pro Gln His Ser Pro Leu Ala Ala Gly Leu Leu
85 90 95
Lys Lys Val Ala Glu Glu Thr Pro Val *
100 105

<210> 1956
<211> 139
<212> PRT
<213> Homo sapiens

<400> 1956
Met Val Leu Pro Phe Ile Cys Asn Leu Leu Arg Arg His Pro Ala Cys
1 5 10 15
Arg Val Leu Val His Arg Pro His Gly Pro Glu Leu Asp Ala Asp Pro
20 25 30
Tyr Asp Pro Gly Glu Glu Asp Pro Ala Gln Ser Arg Ala Leu Glu Ser
35 40 45
Ser Leu Trp Glu Leu Gln Ala Leu Gln Arg His Tyr His Pro Glu Val
50 55 60
Ser Lys Ala Ala Ser Val Ile Asn Gln Ala Leu Ser Met Pro Glu Val
65 70 75 80
Ser Ile Ala Pro Leu Leu Glu Leu Thr Ala Tyr Glu Ile Phe Glu Arg
85 90 95
Asp Leu Lys Lys Lys Gly Pro Glu Pro Val Pro Thr Gly Val Leu Ser
100 105 110
Gln Pro Arg Ala Cys Trp Asp Gly Arg Val Lys Leu Cys Ala Gln His
115 120 125
Phe His Ala Gln Leu Thr Leu Ala His Leu *
130 135 138

<210> 1957
<211> 87
<212> PRT
<213> Homo sapiens

<400> 1957
Met Ala Ala Pro Trp Arg Arg Trp Pro Thr Gly Leu Leu Ala Val Leu
1 5 10 15
Arg Pro Leu Leu Thr Cys Arg Pro Leu Gln Gly Thr Thr Leu Gln Arg
20 25 30
Asp Gly Leu Leu Phe Glu His Asp Arg Gly Arg Phe Phe Thr Ile Leu
35 40 45
Gly Leu Val Cys Ala Gly Gln Gly Gly Phe Trp Ala Ser Met Ala Gly
50 55 60
Ala Gly Ala Leu Arg Thr Pro Gly Pro Leu Gln Gly Met Asn Val Glu
65 70 75 80
Arg His Glu Leu Leu Phe *
85 86

<210> 1958
<211> 48
<212> PRT
<213> Homo sapiens

<400> 1958
Met Thr Tyr Phe Ser Gly Leu Leu Val Ile Leu Ala Phe Ala Ala Trp
1 5 10 15
Val Ala Leu Ala Glu Gly Leu Gly Val Ala Glu Tyr Ala Pro Ala Ala
20 25 30
Leu Pro Cys Ala Ala Cys Ala Thr Ile Leu Leu Ser Ser Val Ala *
35 40 45 47

<210> 1959
<211> 65
<212> PRT
<213> Homo sapiens

<400> 1959
Met Trp Ser Leu Ile Gln Thr Leu Gln Ile Leu Pro Gly Ser Leu Ser
1 5 10 15
Ile Leu Leu Cys Ser Ser Ala Gly Trp Lys Asp Cys Gln Ser Ala Leu
20 25 30
Trp Leu Asn His Val Phe Arg Arg Ala Trp Trp Leu Leu Pro Val Ile
35 40 45
Leu Ala Leu Trp Glu Ala Glu Ala Gly Gly Ser Pro Glu Val Arg Ser
50 55 60 64
*

<210> 1960
<211> 78
<212> PRT
<213> Homo sapiens

<400> 1960

```
Met Ser Tyr Val Arg His Val Leu Ser Cys Leu Gly Gly Gly Leu Ala
 1               5                  10                  15
Leu Trp Arg Ala Gly Gln Trp Leu Trp Ala Gln Arg Leu Gly His Cys
            20                  25                  30
His Thr Tyr Trp Ala Val Ser Glu Glu Leu Leu Pro Asn Ser Gly His
        35                  40                  45
Gly Pro Asp Gly Glu Val Pro Lys Asp Lys Glu Gly Gly Val Phe Asp
    50                  55                  60
Leu Gly Pro Phe Ile Val Gly Phe Trp Gly Pro Gln Ile  *
65                  70                  75      77
```

<210> 1961
<211> 77
<212> PRT
<213> Homo sapiens

<400> 1961

```
Met Trp Tyr Gly Val Phe Leu Trp Ala Leu Val Ser Ser Leu Phe Phe
 1               5                  10                  15
His Val Pro Ala Gly Leu Leu Ala Leu Phe Thr Leu Arg His His Lys
            20                  25                  30
Tyr Gly Ala Ala Ile Ala Gly Val Tyr Arg Ala Ala Gly Lys Glu Met
        35                  40                  45
Ile Pro Phe Glu Ala Leu Thr Leu Gly Thr Gly Gln Thr Phe Cys Val
    50                  55                  60
Leu Val Val Ser Phe Leu Arg Ile Leu Ala Thr Leu  *
65                  70                  75  76
```

<210> 1962
<211> 65
<212> PRT
<213> Homo sapiens

<400> 1962

```
Met Phe Ser Ala Val Phe Pro Ala Val Ser Cys Gln Ile Ser Leu Leu
 1               5                  10                  15
Ser Thr Cys Asn Ser Leu Gln His Phe Pro Tyr Ala Gly Val Leu Cys
            20                  25                  30
Phe Arg Pro Val Leu Cys Leu Cys Pro Gly Gln Asp Phe Cys Gly Asn
        35                  40                  45
Val Arg Cys Gln Trp Arg Leu Leu Ala Gly Val Asp Val Ser Asp Val
    50                  55                  60                  64
 *
```

<210> 1963
<211> 53
<212> PRT
<213> Homo sapiens

<221> misc_feature

<222> (1)...(53)
<223> Xaa = any amino acid or nothing

<400> 1963
Met Thr Cys Pro Leu His Thr Thr Pro Phe Pro Phe Ser Leu Pro Cys
1 5 10 15
Leu Pro Thr Phe Phe Leu Asp Phe Pro Ser Cys Ser Leu Ser Ser Cys
20 25 30
Leu Pro Ile Cys Phe Pro Phe Leu Ser Leu Xaa Gln Ile Leu His Ile
35 40 45
Val Ala Leu Leu Ile
50 53

<210> 1964
<211> 232
<212> PRT
<213> Homo sapiens

<400> 1964
Met Pro Ser Val His Arg Leu Leu Gly Pro Gln Pro Val Pro Ser Arg
1 5 10 15
Arg Leu Arg Leu Ala Leu Ala Leu Leu Leu Ser Leu Gln Val Val Val
20 25 30
Phe Phe Leu Val Val Leu Gly Gln Gly Arg Leu Leu Gln Pro Cys Arg
35 40 45
Gly Cys Leu Glu Leu Pro Gly Gly Pro Gly Glu Ala Glu Asp His Gly
50 55 60
Asp Leu Gly Gln Gly Trp Val Gly Leu Leu Gln Ala Leu Asp Pro Leu
65 70 75 80
Ser His Arg Arg Leu Val Met Ser Thr Arg His Ala His Gly Glu Asp
85 90 95
Arg Ala Phe Leu His Phe Ile Asp Val Lys Leu Val Val Val Pro Ala
100 105 110
Thr Pro His Ile Leu Gln Val Gln Leu His Arg Val Val Glu Val Pro
115 120 125
Leu Leu Arg Arg Leu Phe His Phe Pro Leu Leu Arg Gly Gln Gln Val
130 135 140
Ser Ser Glu Asp Val Val Ile His Thr Leu Val Ala Glu Pro Gln Gly
145 150 155 160
Glu Gly Ala Leu Asn Lys Asp Arg Pro Gly Trp Ile Val Ala Gly Gln
165 170 175
Gly Gly Leu Leu Ile Gly Thr Leu Asp Ser Trp Cys Gly Asp Ile His
180 185 190
Ala Leu Cys Pro Thr Met Trp Gly Trp Gly Gly Ser Ala Ala Pro Val
195 200 205
Glu Ser Leu Gly Lys Gly Thr Ser Gly Glu Gly Asp Gly Arg Arg Gln
210 215 220
Gly Gln Arg Thr Gly Pro Gly *
225 230 231

<210> 1965
<211> 253
<212> PRT

<213> Homo sapiens

<400> 1965
Met Gly Cys Ala Ile Ile Ala Gly Phe Leu His Tyr Leu Phe Leu Ala
1 5 10 15
Cys Phe Phe Trp Met Leu Val Glu Ala Val Ile Leu Phe Leu Met Val
20 25 30
Arg Asn Leu Lys Val Val Asn Tyr Phe Ser Ser Arg Asn Ile Lys Met
35 40 45
Leu His Ile Cys Ala Phe Gly Tyr Gly Leu Pro Met Leu Val Val Val
50 55 60
Ile Ser Ala Ser Val Gln Pro Gln Gly Tyr Gly Met His Asn Arg Cys
65 70 75 80
Trp Leu Asn Thr Glu Thr Gly Phe Ile Trp Ser Phe Leu Gly Pro Val
85 90 95
Cys Thr Val Ile Val Ile Asn Ser Leu Leu Leu Thr Trp Thr Leu Trp
100 105 110
Ile Leu Arg Gln Arg Leu Ser Ser Val Asn Ala Glu Val Ser Thr Leu
115 120 125
Lys Asp Thr Arg Leu Leu Thr Phe Lys Ala Phe Ala Gln Leu Phe Ile
130 135 140
Leu Gly Cys Ser Trp Val Leu Gly Ile Phe Gln Ile Gly Pro Val Ala
145 150 155 160
Gly Val Met Ala Tyr Leu Phe His His His Gln Gln Pro Ala Gly Gly
165 170 175
Leu His Leu Pro His Pro Leu Ser Ala Gln Arg Pro Gly Thr Arg Arg
180 185 190
Ile Gln Glu Val Asp His Trp Glu Asp Glu Ala Gln Leu Pro Val Pro
195 200 205
Asp Leu Lys Asp Leu Ala Val Leu His Ala Ile Arg Phe Gln Asp Gly
210 215 220
Leu Lys Ser Phe Leu Ala Phe Lys Tyr Ala Met Glu Pro Thr Val Gly
225 230 235 240
Gly Thr Ser Ser Phe Pro Cys Arg Glu Pro Tyr Pro *
245 250 252

<210> 1966
<211> 649
<212> PRT
<213> Homo sapiens

<400> 1966
Met Val Thr Cys Phe Ile Ile Gly Leu Leu Phe Pro Val Phe Ser Val
1 5 10 15
Cys Tyr Leu Ile Ala Pro Lys Ser Pro Leu Gly Leu Phe Ile Arg Lys
20 25 30
Pro Phe Ile Lys Phe Ile Cys His Thr Ala Ser Tyr Leu Thr Phe Leu
35 40 45
Phe Leu Leu Leu Leu Ala Ser Gln His Ile Asp Arg Ser Asp Leu Asn
50 55 60
Arg Gln Gly Pro Pro Pro Thr Ile Val Glu Trp Met Ile Leu Pro Trp
65 70 75 80
Val Leu Gly Phe Ile Trp Gly Glu Ile Lys Gln Met Trp Asp Gly Gly
85 90 95
Leu Gln Asp Tyr Ile His Asp Trp Trp Asn Leu Met Asp Phe Val Met

```
            100                 105                 110
Asn Ser Leu Tyr Leu Ala Thr Ile Ser Leu Lys Ile Val Ala Phe Val
        115                 120                 125
Lys Tyr Ser Ala Leu Asn Pro Arg Glu Ser Trp Asp Met Trp His Pro
    130                 135                 140
Thr Leu Val Ala Glu Ala Leu Phe Ala Ile Ala Asn Ile Phe Ser Ser
145                 150                 155                 160
Leu Arg Leu Ile Ser Leu Phe Thr Ala Asn Ser His Leu Gly Pro Leu
                165                 170                 175
Gln Ile Ser Leu Gly Arg Met Leu Leu Asp Ile Leu Lys Phe Leu Phe
            180                 185                 190
Ile Tyr Cys Leu Val Leu Leu Ala Phe Ala Asn Gly Leu Asn Gln Leu
        195                 200                 205
Tyr Phe Tyr Tyr Glu Glu Thr Lys Gly Leu Thr Cys Lys Gly Ile Arg
    210                 215                 220
Cys Glu Lys Gln Asn Asn Ala Phe Ser Thr Leu Phe Glu Thr Leu Gln
225                 230                 235                 240
Ser Leu Phe Trp Ser Ile Phe Gly Leu Ile Asn Leu Tyr Val Thr Asn
                245                 250                 255
Val Lys Ala Gln His Glu Phe Thr Glu Phe Val Gly Ala Thr Met Phe
            260                 265                 270
Gly Thr Tyr Asn Asp Ile Ser Leu Val Val Leu Leu Asn Met Leu Ile
        275                 280                 285
Ala Met Met Asn Asn Ser Tyr Gln Leu Ile Ala Asp His Ala Asp Ile
    290                 295                 300
Glu Trp Lys Phe Ala Arg Thr Lys Leu Trp Met Ser Tyr Phe Glu Glu
305                 310                 315                 320
Gly Gly Thr Leu Pro Thr Pro Phe Asn Val Ile Pro Ser Pro Lys Ser
                325                 330                 335
Leu Trp Tyr Leu Ile Lys Trp Ile Trp Thr His Leu Cys Lys Lys Lys
            340                 345                 350
Met Arg Arg Lys Pro Glu Ser Phe Gly Thr Ile Gly Arg Arg Ala Ala
        355                 360                 365
Asp Asn Leu Arg Arg His His Gln Tyr Gln Glu Val Met Arg Asn Leu
    370                 375                 380
Val Lys Arg Tyr Val Ala Ala Met Ile Arg Asp Ala Lys Thr Glu Glu
385                 390                 395                 400
Gly Leu Thr Glu Glu Asn Phe Lys Glu Leu Lys Gln Asp Ile Ser Ser
                405                 410                 415
Phe Arg Phe Glu Val Leu Gly Leu Leu Arg Gly Ser Lys Leu Ser Thr
            420                 425                 430
Ile Gln Ser Ala Asn Ala Ser Lys Glu Ser Ser Asn Ser Ala Asp Ser
        435                 440                 445
Asp Glu Lys Ser Asp Ser Glu Gly Asn Ser Lys Asp Lys Lys Lys Asn
    450                 455                 460
Phe Ser Leu Phe Asp Leu Thr Thr Leu Ile His Pro Arg Ser Ala Ala
465                 470                 475                 480
Ile Ala Ser Glu Arg His Asn Ile Ser Asn Gly Ser Ala Leu Val Val
                485                 490                 495
Gln Glu Pro Pro Arg Glu Lys Gln Arg Lys Val Asn Phe Val Thr Asp
            500                 505                 510
Ile Lys Asn Phe Gly Leu Phe His Arg Arg Ser Lys Gln Asn Ala Ala
        515                 520                 525
Glu Gln Asn Ala Asn Gln Ile Phe Ser Val Ser Glu Glu Val Ala Arg
    530                 535                 540
Gln Gln Ala Ala Gly Pro Leu Glu Arg Asn Ile Gln Leu Glu Ser Arg
545                 550                 555                 560
Gly Leu Ala Ser Arg Gly Asp Leu Ser Ile Pro Gly Leu Ser Glu Gln
                565                 570                 575
```

```
Cys Val Leu Val Asp His Arg Glu Arg Asn Thr Asp Thr Leu Gly Leu
            580                 585                 590
Gln Val Gly Lys Arg Val Cys Pro Phe Lys Ser Glu Lys Val Val Val
        595                 600                 605
Glu Asp Thr Val Pro Ile Ile Pro Lys Glu Lys His Ala Lys Glu Glu
    610                 615                 620
Asp Ser Ser Ile Asp Tyr Asp Leu Asn Leu Pro Asp Thr Val Thr His
625                 630                 635                 640
Glu Asp Tyr Val Thr Thr Arg Leu  *
                645         648
```

<210> 1967
<211> 80
<212> PRT
<213> Homo sapiens

<400> 1967

```
Met Thr Gly Thr His Gln Tyr Ala Trp Val Ile Phe Val Phe Leu Ser
 1               5                  10                  15
Thr Tyr Arg Ile Ser Pro Cys Trp Pro Gly Trp Phe Gln Thr Pro Gly
            20                  25                  30
Leu Arg Trp Ser Ala Cys Leu Gly Leu Pro Gly Cys Trp Asp Cys Arg
        35                  40                  45
Arg Glu Pro Leu Gly Pro Ala Cys Ile Phe Tyr Gln Pro Gln Ile Gln
    50                  55                  60
Gln Gln Ala Glu Asp Ser Ala His Lys Thr Gly Leu Val Ser Trp  *
65                  70                  75                  79
```

<210> 1968
<211> 49
<212> PRT
<213> Homo sapiens

<400> 1968

```
Met Thr Tyr Ile Leu Val Tyr Lys Leu Gly Ser Ile Leu Leu Ser Phe
 1               5                  10                  15
Phe Leu Ile Cys Phe Glu Glu Phe Ser Ser Glu Asn Ser Gly Pro Gly
            20                  25                  30
Ile Phe Phe Val Glu Arg Val Leu Ile Leu Asn Leu Ile Ser Leu Ile
        35                  40                  45          48
 *
```

<210> 1969
<211> 150
<212> PRT
<213> Homo sapiens

<400> 1969

```
Met His Val His Phe Trp Leu Val Thr Ala Ser Phe Ser Ser Ser Val
```

```
 1               5                   10                  15
Ala Trp Thr Thr Ala Glu Ile Thr Gly Gly Val Ser Gly Val Ala Ala
            20                  25                  30
Gly Val Gly Ser Trp Glu Gly Gly Ser Glu Arg Gly Asp Arg Phe Gly
        35                  40                  45
Asp Phe Phe Thr Leu Asn Val Ser Val Phe Arg Gly Val Phe Phe Phe
    50                  55                  60
Leu Ala Gly Leu Phe Ser Pro Ser Pro Ser Thr Pro Leu Ala Ser Ile
65                  70                  75                  80
Ala Leu Ala Gly Ile Ser Lys Glu Ala Gly Asp Leu Glu Gly Glu Leu
                85                  90                  95
Gly Val Leu Glu Asp Val Leu Lys Gly Ser Thr Asp Ser Ser Gln Val
            100                 105                 110
Ser Gly Ser Lys Leu Tyr Asp Cys Trp Gly Ser Leu Gly Asp Ser Cys
        115                 120                 125
Ile Phe Glu Val Glu Glu Lys Gly Leu Lys Leu Gly Ser Ser His Leu
    130                 135                 140
Ser Ile Ser Lys Val  *
145             149
```

<210> 1970
<211> 48
<212> PRT
<213> Homo sapiens

<400> 1970

```
Met Phe Gly Ser Arg Gly Leu Leu Cys Met Cys Val Phe Phe Phe Asn
 1               5                   10                  15
Ile Leu Ala Ser Gln Cys Lys Val Ile Ser Ser Gly Gly Met Leu Cys
            20                  25                  30
Cys Arg Thr Pro Thr Leu Leu Asp Tyr Leu Arg Gln His Phe Leu  *
        35                  40                  45      47
```

<210> 1971
<211> 64
<212> PRT
<213> Homo sapiens

<400> 1971

```
Met Leu Ile Phe Thr Val Leu Glu Leu Leu Leu Ala Ala Tyr Ser Ser
 1               5                   10                  15
Val Phe Trp Trp Lys Gln Leu Tyr Ser Asn Asn Pro Gly Val Ser Met
            20                  25                  30
Leu Thr Cys Arg Leu Ile Pro Ala Val Ser Gln Val Gln Ala Thr Ile
        35                  40                  45
Ile Gln Pro Gln Lys Val Ala Lys Arg Arg Ile Asn Tyr Cys Ser  *
    50                  55                  60          63
```

<210> 1972
<211> 211
<212> PRT

<213> Homo sapiens

<221> misc_feature
<222> (1)...(211)
<223> Xaa = any amino acid or nothing

<400> 1972
Met Thr Arg Met Leu Asn Met Leu Ile Val Phe Arg Phe Leu Arg Ile
1 5 10 15
Ile Pro Ser Met Lys Pro Met Ala Val Val Ala Ser Thr Val Leu Gly
20 25 30
Leu Val Gln Asn Met Arg Ala Phe Gly Gly Ile Leu Val Val Val Tyr
35 40 45
Tyr Val Phe Ala Ile Ile Gly Ile Asn Leu Phe Arg Gly Val Ile Val
50 55 60
Ala Leu Pro Gly Asn Ser Ser Leu Ala Pro Ala Asn Gly Ser Ala Pro
65 70 75 80
Cys Gly Ser Phe Glu Gln Leu Glu Tyr Trp Ala Asn Asn Phe Asp Asp
85 90 95
Phe Xaa Ala Ala Leu Val Thr Leu Trp Asn Leu Met Val Val Asn Asn
100 105 110
Trp Gln Val Phe Leu Asp Ala Tyr Arg Arg Tyr Ser Gly Pro Trp Ser
115 120 125
Lys Ile Tyr Phe Val Leu Trp Trp Leu Val Ser Ser Val Ile Trp Val
130 135 140
Asn Leu Phe Leu Ala Leu Ile Leu Glu Asn Phe Leu His Lys Trp Asp
145 150 155 160
Pro Arg Ser His Leu Gln Pro Leu Ala Gly Thr Pro Glu Ala Thr Tyr
165 170 175
Gln Met Thr Val Glu Leu Leu Phe Arg Asp Ile Leu Glu Glu Pro Gly
180 185 190
Glu Asp Glu Leu Thr Glu Arg Leu Ser Gln His Pro His Leu Trp Leu
195 200 205
Cys Arg *
210

<210> 1973
<211> 53
<212> PRT
<213> Homo sapiens

<400> 1973
Met Ile Gln Tyr Ala Val Phe Val Leu Cys Gly Phe Leu Tyr Leu Cys
1 5 10 15
Phe Met Leu Phe Phe Phe Ser Ser Val Thr Gln Ala Gly Val Ser Glu
20 25 30
Pro Arg Ser Ser His Cys Thr Pro Ala Trp Ala Thr Glu Arg Asp Cys
35 40 45
Val Ser Asn Lys *
50 52

<210> 1974
<211> 50

<212> PRT
<213> Homo sapiens

<400> 1974
Met Gly Val Thr Thr Ala Thr Leu Ile Ala Pro Ala Leu Arg Thr Leu
1 5 10 15
Arg Thr Ser Ala Val Cys Ser Thr Thr Ala Glu Thr Ser Phe Ser Ala
20 25 30
Cys Thr Phe Val Ser Thr Ser Cys Ser Lys Lys Gly Thr Pro Arg Phe
35 40 45
Ser *
49

<210> 1975
<211> 87
<212> PRT
<213> Homo sapiens

<400> 1975
Met Cys Ser Ser Pro Ala Val Leu Leu Cys Ala Leu Val Val Gly Cys
1 5 10 15
Pro Val Gly Phe Pro His Glu Ala Asp Pro Gly Ser Met Gln Arg Ala
20 25 30
Ser Ser Leu Gly Leu His Gln Ala Ser Val Val Ser Ala Gly Trp Leu
35 40 45
Gly Gln Ala Arg His Gly Ala His Leu Gly Cys Ser Leu Leu Pro Ser
50 55 60
Gly Val His Gly Leu Trp Arg Pro Ser Val Gln Pro Arg Arg Asp Pro
65 70 75 80
Val Thr Glu Leu Gln Cys *
85 86

<210> 1976
<211> 107
<212> PRT
<213> Homo sapiens

<400> 1976
Met Ala Leu Tyr Glu Leu Phe Ser His Pro Val Glu Arg Ser Tyr Arg
1 5 10 15
Ala Gly Leu Cys Ser Lys Ala Ala Leu Phe Leu Leu Leu Ala Ala Ala
20 25 30
Leu Thr Tyr Ile Pro Pro Leu Leu Val Ala Phe Arg Ser His Gly Phe
35 40 45
Trp Leu Lys Arg Ser Ser Tyr Glu Glu Gln Pro Thr Val Arg Phe Gln
50 55 60
His Gln Val Leu Leu Val Ala Leu Leu Gly Pro Glu Ser Asp Gly Phe
65 70 75 80
Leu Ala Trp Ser Thr Phe Pro Ala Phe Asn Arg Gln Gln Gly Asp Arg
85 90 95
Leu Arg Val Pro Leu Val Ser Trp Arg Arg *
100 105 106

<210> 1977
<211> 134
<212> PRT
<213> Homo sapiens

<400> 1977
Met Val Thr Val Ala Met Ala Cys Ser Gly Ala Leu Thr Ala Leu Cys
1 5 10 15
Cys Leu Phe Val Ala Met Gly Val Leu Arg Val Pro Trp His Cys Pro
20 25 30
Leu Leu Leu Val Thr Glu Gly Leu Leu Asp Met Leu Ile Ala Gly Gly
35 40 45
Tyr Ile Pro Ala Leu Tyr Phe Tyr Phe His Tyr Leu Ser Ala Ala Tyr
50 55 60
Gly Ser Pro Val Cys Lys Glu Arg Gln Ala Leu Tyr Gln Ser Lys Gly
65 70 75 80
Tyr Ser Gly Phe Gly Cys Ser Phe His Gly Ala Asp Ile Gly Ala Gly
85 90 95
Ile Phe Ala Ala Leu Gly Ile Val Val Phe Ala Leu Gly Ala Val Leu
100 105 110
Ala Ile Lys Gly Tyr Arg Lys Val Arg Lys Leu Lys Glu Lys Pro Ala
115 120 125
Glu Met Phe Glu Phe *
130 133

<210> 1978
<211> 61
<212> PRT
<213> Homo sapiens

<400> 1978
Met Thr Leu Arg Met Leu Val Pro Arg Leu Leu Leu Thr Arg Gln Leu
1 5 10 15
Val Trp Phe Phe Ser Ala Ala Thr Glu Arg Asp Pro Glu Met Met Asn
20 25 30
Gly Ile Pro Arg Lys Leu Met Ser Phe Pro Pro Ser Ser Val Thr Ser
35 40 45
Arg Arg Ser Arg Arg Gly His His Leu Gln Ser Leu *
50 55 60

<210> 1979
<211> 66
<212> PRT
<213> Homo sapiens

<400> 1979
Met Leu Thr Ala Leu Pro Lys Ser Phe Val Phe Lys Val Val Gly Glu
1 5 10 15
Trp Trp Trp Leu Phe Ile Cys Leu Val Leu Ala Phe Ala Asp Gly Lys

```
                20                  25                  30
Arg His Lys Tyr Ser Tyr Asp Ala Asn Val Phe Leu Gln Val Asn Tyr
            35                  40                  45
Ile Thr Trp Pro Asp Ser Phe Ser Pro Val Pro Ser Leu Pro Pro Ile
        50                  55                  60
Leu  *
 65
```

<210> 1980
<211> 51
<212> PRT
<213> Homo sapiens

<400> 1980

```
Met Asp Thr Pro Arg Ser Thr Val Phe Ser Leu Trp Phe Gly Ile His
  1               5                  10                  15
Lys Ala Ala Gly Ile Phe Gln Val Leu Val Gln Leu Leu Leu Leu Leu
                20                  25                  30
Thr Pro Tyr Pro Arg Tyr Pro Ser Pro Ser Pro Leu Pro Pro Tyr Ser
            35                  40                  45
Tyr Pro  *
     50
```

<210> 1981
<211> 79
<212> PRT
<213> Homo sapiens

<400> 1981

```
Met Met Trp Ala Ala Gly Ala Val Ala Ala Met Ser Ser Ile Thr Phe
  1               5                  10                  15
Pro Ala Val Ser Ala Leu Val Ser Arg Thr Ala Asp Ala Asp Gln Gln
                20                  25                  30
Gly Glu Leu Ile Gly Thr Ser Asp Asn Tyr Leu Lys Val Gln Asn Val
            35                  40                  45
Leu Ile Leu Cys Ser Val Ser Tyr Val Leu Lys His Lys Tyr Ile Phe
        50                  55                  60
Arg Gly Glu Thr Phe Lys Ile Ala Phe Asp Ile Asn Arg Lys Ser
 65                  70                  75                  79
```

<210> 1982
<211> 156
<212> PRT
<213> Homo sapiens

<400> 1982

```
Met His Asn Asn Tyr Thr Ala Leu Leu Gly Val Trp Ile Tyr Gly Phe
  1               5                  10                  15
Phe Val Leu Met Leu Leu Val Leu Asp Leu Leu Tyr Tyr Ser Ala Met
                20                  25                  30
```

Asn Tyr Asp Ile Cys Lys Val Tyr Leu Ala Arg Trp Gly Ile Gln Gly
35 40 45
Arg Trp Met Lys Gln Asp Pro Arg Arg Trp Gly Asn Pro Ala Arg Ala
50 55 60
Pro Arg Pro Gly Gln Arg Ala Pro Gln Pro Gln Pro Pro Pro Gly Pro
65 70 75 80
Leu Pro Gln Ala Pro Gln Ala Val His Thr Leu Arg Gly Asp Ala His
85 90 95
Ser Pro Pro Leu Met Thr Phe Gln Ser Ser Ser Ala Trp Glu Gly Ala
100 105 110
Ser Gln Gln Gln Glu Ile Pro Glu Asn Glu Glu Thr Glu Lys Gly Asp
115 120 125
Asp Gln Ile Ser Ser Phe Leu Gly Val Thr Ser Asn Thr Lys Glu Ala
130 135 140
Ser Val Ile Gly Ile Gln Lys Thr Val Asp Val Leu
145 150 155 156

<210> 1983
<211> 63
<212> PRT
<213> Homo sapiens

<400> 1983
Met Arg Leu Ile Arg Ile Trp Phe Ser Gly Lys Phe Phe Pro Ala Gly
1 5 10 15
Leu His Ser Gln Ser Leu Pro Ser Ile Ser Ala Ala Ile Gly Leu Leu
20 25 30
Met Leu Phe Thr Asn Leu Phe Thr Cys Ser Lys Cys Phe Val Ile Ser
35 40 45
Val Ala Lys Thr Met Ser Ile Ile Ala Trp Arg Ser Val Arg *
50 55 60 62

<210> 1984
<211> 232
<212> PRT
<213> Homo sapiens

<400> 1984
Met Phe His Arg Cys Gly Ile Met Ala Leu Val Ala Ala Tyr Leu Asn
1 5 10 15
Phe Val Ser Gln Met Ile Ala Val Pro Ala Phe Cys Gln His Val Ser
20 25 30
Lys Val Ile Glu Ile Arg Thr Met Glu Ala Pro Tyr Phe Leu Pro Glu
35 40 45
His Ile Phe Arg Asp Lys Cys Met Leu Pro Lys Ser Leu Glu Lys His
50 55 60
Glu Lys Asp Leu Tyr Phe Leu Thr Asn Lys Ile Ala Glu Ser Leu Gly
65 70 75 80
Gly Lys Trp Asp Ile Val Leu Arg Asp Cys Gln Phe Arg Met Leu Pro
85 90 95
Gln Val Thr Asp Glu Asp Arg Leu Ser Arg Arg Lys Ser Ile Val Asp
100 105 110
Thr Val Ser Ile Gln Val Asp Ile Leu Ser Asn Asn Val Pro Ser Asp

115 120 125
Asp Val Val Ser Asn Thr Glu Glu Ile Thr Phe Glu Ala Leu Lys Lys
130 135 140
Ala Ile Asp Thr Ser Gly Met Glu Glu Gln Glu Lys Glu Lys Arg Arg
145 150 155 160
Leu Val Ile Glu Lys Phe Gln Lys Ala Pro Phe Glu Glu Ile Ala Ala
165 170 175
Gln Cys Glu Ser Lys Ala Asn Leu Leu His Asp Arg Leu Ala Gln Ile
180 185 190
Leu Glu Leu Thr Ile Arg Pro Pro Pro Ser Pro Ser Gly Thr Leu Thr
195 200 205
Ile Thr Ser Gly His Ala Gln Tyr Gln Ser Val Pro Val Tyr Glu Met
210 215 220
Lys Phe Pro Asp Leu Cys Val Tyr
225 230 232

<210> 1985
<211> 141
<212> PRT
<213> Homo sapiens

<400> 1985
Met Asn Leu Ser Leu Pro Phe Leu Cys Leu Phe Leu Leu Ser Phe Ser
1 5 10 15
Phe Lys Leu Ala Leu Gln Leu Arg Lys Val Ser Leu Leu Ser Leu Arg
20 25 30
Leu Trp Gly Gln Ser Ile Cys Cys Leu Glu Lys Glu Gly Asn Gln Asp
35 40 45
Ser Ser Gly Thr Gln Met Ser Ser Ser Leu Ala Leu Leu Asn Pro Leu
50 55 60
Leu His Asn Trp Ser Phe Ile Leu Ala Leu Asn Asp Pro Ala Gly His
65 70 75 80
His Gly Phe Leu Phe Leu Leu Val Phe Phe Phe Ser Glu Thr Glu Ser
85 90 95
His Ser Val Thr Gln Ala Gly Val Gln Trp Arg Asp Leu Ser Ser Leu
100 105 110
Gln Pro Leu Pro Pro Gly Phe Lys Arg Phe Phe Cys Leu Ser Leu Pro
115 120 125
Ser Ser Trp Asp Tyr Arg Cys Ala Thr Thr Pro Gly *
130 135 140

<210> 1986
<211> 292
<212> PRT
<213> Homo sapiens

<400> 1986
Met Ile Ser Val Ser Ala Met Ala Ile Ala Phe Leu Thr Leu Gly Tyr
1 5 10 15
Phe Phe Lys Ile Lys Glu Ile Lys Ser Pro Glu Met Ala Glu Asp Trp
20 25 30
Asn Thr Phe Leu Leu Arg Phe Asn Asp Leu Asp Leu Cys Val Ser Glu
35 40 45

```
Asn Glu Thr Leu Lys His Leu Thr Asn Asp Thr Thr Thr Pro Glu Ser
     50                  55                  60
Thr Met Thr Ser Gly Gln Ala Arg Ala Ser Thr Gln Ser Pro Gln Ala
 65                  70                  75                  80
Leu Glu Asp Ser Gly Pro Val Asn Ile Ser Val Ser Ile Thr Leu Thr
                 85                  90                  95
Leu Asp Pro Leu Lys Pro Phe Gly Gly Tyr Ser Arg Asn Val Thr His
            100                 105                 110
Leu Tyr Ser Thr Ile Leu Gly His Gln Ile Gly Leu Ser Gly Arg Glu
        115                 120                 125
Ala His Glu Glu Ile Asn Ile Thr Phe Thr Leu Pro Thr Ala Trp Ser
    130                 135                 140
Ser Asp Asp Cys Ala Leu His Gly His Cys Glu Gln Val Val Phe Thr
145                 150                 155                 160
Ala Cys Met Thr Leu Thr Ala Ser Pro Gly Val Phe Pro Val Thr Val
                165                 170                 175
Gln Pro Pro His Cys Val Pro Asp Thr Tyr Ser Asn Ala Thr Leu Trp
            180                 185                 190
Tyr Lys Ile Phe Thr Thr Ala Arg Asp Ala Asn Thr Lys Tyr Ala Gln
        195                 200                 205
Asp Tyr Asn Pro Phe Trp Cys Tyr Lys Gly Ala Ile Gly Lys Val Tyr
    210                 215                 220
His Ala Leu Asn Pro Lys Leu Thr Val Ile Val Pro Asp Asp Asp Arg
225                 230                 235                 240
Ser Leu Ile Asn Leu His Leu Met His Thr Ser Tyr Phe Leu Phe Val
                245                 250                 255
Met Val Ile Thr Met Phe Cys Tyr Ala Val Ile Lys Gly Arg Pro Ser
            260                 265                 270
Lys Leu Arg Gln Ser Asn Pro Glu Phe Cys Pro Glu Lys Val Ala Leu
        275                 280                 285
Ala Glu Ala  *
    290 291
```

<210> 1987
<211> 186
<212> PRT
<213> Homo sapiens

<400> 1987

```
Met Ala Gly Pro Arg Pro Arg Trp Arg Asp Gln Leu Leu Phe Met Ser
  1               5                  10                  15
Ile Ile Val Leu Val Ile Val Val Ile Cys Leu Met Leu Tyr Ala Leu
             20                  25                  30
Leu Trp Glu Ala Gly Asn Leu Thr Asp Leu Pro Asn Leu Arg Ile Gly
         35                  40                  45
Phe Tyr Asn Phe Cys Leu Trp Asn Glu Asp Thr Ser Thr Leu Gln Cys
     50                  55                  60
His Gln Phe Pro Glu Leu Glu Ala Leu Gly Val Pro Arg Val Gly Leu
 65                  70                  75                  80
Gly Leu Ala Arg Leu Gly Val Tyr Gly Ser Leu Val Leu Thr Leu Phe
                 85                  90                  95
Ala Pro Gln Pro Leu Leu Leu Ala Gln Cys Asn Ser Asp Glu Arg Ala
            100                 105                 110
Trp Arg Leu Ala Val Gly Phe Leu Ala Val Ser Ser Val Leu Leu Ala
        115                 120                 125
Gly Gly Leu Gly Leu Phe Leu Ser Tyr Val Trp Lys Trp Val Arg Leu
```

```
    130                 135                 140
Ser Leu Pro Gly Pro Gly Phe Leu Ala Leu Gly Ser Ala Gln Ala Leu
145                 150                 155                 160
Leu Ile Leu Leu Leu Ile Ala Met Ala Val Phe Pro Leu Arg Ala Glu
                165                 170                 175
Arg Ala Glu Ser Lys Leu Glu Ser Cys  *
            180                 185
```

<210> 1988
<211> 47
<212> PRT
<213> Homo sapiens

<400> 1988

```
Met Phe Asn Leu Lys Glu Ile Pro Leu Ile Leu Tyr Val Leu Leu Ser
 1               5                  10                  15
Val Val Cys Phe Ser Phe Ser Tyr Gly Val Glu Pro Pro Lys Ser Trp
            20                  25                  30
Ser Gln Gly Lys Lys Gly Val Val Thr Gly Asp Ser Leu Leu  *
        35                  40                  45  46
```

<210> 1989
<211> 58
<212> PRT
<213> Homo sapiens

<400> 1989

```
Met Thr Leu Pro Cys Ala Ile Gln Met Phe Ile Ala Ala Val Gln Val
 1               5                  10                  15
Leu Ser Val Thr Tyr Leu Asp Leu Gln Pro His Leu Asn Glu Ser Leu
            20                  25                  30
Leu Thr Val Ser Leu Ile Phe Arg Phe Ile Phe Asn Leu Leu Phe Tyr
        35                  40                  45
Leu Gly Leu Thr Phe Ser Val Thr Lys  *
    50                  55      57
```

<210> 1990
<211> 80
<212> PRT
<213> Homo sapiens

<400> 1990

```
Met Ile Ser Phe Val Leu Val Lys Gly Leu Phe Leu Lys Cys Thr Phe
 1               5                  10                  15
His Phe Pro Leu Phe Asn Arg His Ile Met Ser Cys Ser Phe Leu Arg
            20                  25                  30
Ser Asp Phe Met His Gly Asp Ser Met Cys Phe Ser Ser Ser Tyr Met
        35                  40                  45
Leu Leu Asn Glu Ser Leu Tyr Ile Ser Phe His Thr Met Val Ile Lys
    50                  55                  60
```

Thr His Trp Ala Val Cys Gly Cys Gly Phe Ile Ser Glu Lys Leu *
65 70 75 79

<210> 1991
<211> 48
<212> PRT
<213> Homo sapiens

<400> 1991
Met Val Arg Trp Lys Arg Glu Ile His Glu Leu Leu Trp Pro Leu Trp
1 5 10 15
Phe Cys Ser Trp Pro Arg Val Phe Glu Lys Gln Arg Ser Met Thr Asp
20 25 30
Phe Thr Cys Ser Ala Phe Ser Ala Phe Cys Leu Phe Cys Cys Pro *
35 40 45 47

<210> 1992
<211> 51
<212> PRT
<213> Homo sapiens

<400> 1992
Met Leu Phe Ser Leu Gln Thr Ala Ile Val Tyr Cys Thr Ile Thr Val
1 5 10 15
Leu Cys His Arg Thr Leu Ile Phe Ser Ser Met His Lys Cys Ile Met
20 25 30
Leu Phe Pro Ile Ile His Ile Cys Ser Tyr Val Phe Phe Val Ile Tyr
35 40 45
Ser Phe *
50

<210> 1993
<211> 79
<212> PRT
<213> Homo sapiens

<400> 1993
Met Trp Cys Ala Glu Met Leu His Ile Leu Phe Met Gly Leu Arg Val
1 5 10 15
Asn Leu Asn His Glu Thr Phe Leu Ile Ile Cys Cys Glu Ile Tyr Gln
20 25 30
Ala Trp Met Ile Ser Val Phe Leu Val Val Cys Cys Phe Phe Lys Glu
35 40 45
Val Ile Gln Val Pro Leu Leu Ser Cys Gln His Thr Lys Leu Leu Lys
50 55 60
Lys Leu Thr Ile Ser Phe Arg Ser Asn Ser Gln Pro Val Glu *
65 70 75 78

<210> 1994
<211> 52
<212> PRT
<213> Homo sapiens

<400> 1994
Met Thr Ser Leu Gln Lys Arg Leu Leu Ser His Cys Met Gln Cys Thr
1 5 10 15
Met Leu Leu Gly Ile Cys Gly Gln Cys Lys Asp Asp Asp Ile Leu Ala
20 25 30
Ser Trp Val Ile Gln Glu Phe Thr Ala Met Gln Ser Arg Ser Arg Asn
35 40 45
Leu Gln Ser Arg
50 52

<210> 1995
<211> 164
<212> PRT
<213> Homo sapiens

<400> 1995
Met Leu Leu Ala Thr Leu Leu Leu Leu Leu Leu Gly Gly Ala Leu Ala
1 5 10 15
His Pro Asp Arg Ile Ile Phe Pro Asn His Ala Cys Glu Asp Pro Pro
20 25 30
Ala Val Leu Leu Glu Val Gln Gly Thr Leu Gln Arg Pro Leu Val Arg
35 40 45
Asp Ser Arg Thr Ser Pro Ala Asn Cys Thr Trp Leu Ile Leu Gly Ser
50 55 60
Lys Glu Arg Thr Val Thr Ile Arg Phe Gln Lys Leu His Leu Ala Cys
65 70 75 80
Gly Ser Glu Arg Leu Thr Leu Arg Ser Pro Leu Gln Pro Leu Ile Ser
85 90 95
Leu Cys Glu Ala Pro Pro Ser Pro Leu Gln Leu Pro Gly Gly Asn Val
100 105 110
Thr Ile Thr Tyr Ser Tyr Ala Gly Gly Gln Ser Thr His Gly Pro Gly
115 120 125
Leu Pro Ala Leu Leu Gln Ala Ser Pro Ser Pro Trp Cys Leu Cys Arg
130 135 140
Leu Ala Asp Val Leu Ala Arg Arg Gly Ser Met Pro Glu Pro Pro Leu
145 150 155 160
Cys Ile Cys *
163

<210> 1996
<211> 77
<212> PRT
<213> Homo sapiens

<400> 1996
Met Trp Tyr Gly Val Phe Leu Trp Ala Leu Val Ser Ser Leu Phe Phe
1 5 10 15

His Val Pro Ala Gly Leu Leu Ala Leu Phe Thr Leu Arg His His Lys
20 25 30
Tyr Gly Ala Ala Ile Ala Gly Val Tyr Arg Ala Ala Gly Lys Glu Met
35 40 45
Ile Pro Phe Glu Ala Leu Thr Leu Gly Thr Gly Gln Thr Phe Cys Val
50 55 60
Leu Val Val Ser Phe Leu Arg Ile Leu Ala Thr Leu *
65 70 75 76

<210> 1997
<211> 233
<212> PRT
<213> Homo sapiens

<400> 1997
Met Gly Leu Pro Gly Leu Phe Cys Leu Ala Val Leu Ala Ala Ser Ser
1 5 10 15
Phe Ser Lys Ala Arg Glu Glu Glu Ile Thr Pro Val Val Ser Ile Ala
20 25 30
Tyr Lys Val Leu Glu Val Phe Pro Lys Gly Arg Trp Val Leu Ile Thr
35 40 45
Cys Cys Ala Pro Gln Pro Pro Pro Pro Ile Thr Tyr Ser Leu Cys Gly
50 55 60
Thr Lys Asn Ile Lys Val Ala Lys Lys Val Val Lys Thr His Glu Pro
65 70 75 80
Ala Ser Phe Asn Leu Asn Val Thr Leu Lys Ser Ser Pro Asp Leu Leu
85 90 95
Thr Tyr Phe Cys Arg Ala Ser Ser Thr Ser Gly Ala His Val Asp Ser
100 105 110
Ala Arg Leu Gln Met His Trp Glu Leu Trp Ser Arg Gln Arg Gly Arg
115 120 125
Pro Gln Gly Gly Asp Asp Leu Pro Gly Val Leu Gly Gln Pro Thr Tyr
130 135 140
His Gln Gln Pro Asp Arg Glu Gly Trp Ala Gly Pro Pro Ala Ala Glu
145 150 155 160
Thr Met Pro Gln Glu Ala Cys Gln Leu Ser Pro Ser Cys Arg Ala Arg
165 170 175
His Arg Thr Trp Phe Trp Cys Gln Ala Cys Lys Gln Arg Gln Cys Ser
180 185 190
Ser Thr Ala Pro Ser Gln Trp Leu Pro Gln Val Val Thr Gln Lys Met
195 200 205
Glu Asp Trp Gln Gly Pro Pro Gly Glu Pro His Pro Cys Leu Ala Ala
210 215 220
Leu Gln Glu His Pro Pro Ser Glu *
225 230 232

<210> 1998
<211> 58
<212> PRT
<213> Homo sapiens

<400> 1998
Met Pro Ala Ile Val Val Phe Leu Phe Cys Phe Val Ile Ser Asp Gly

```
 1               5                   10                  15
Leu Thr Leu Ser Pro Arg Leu Asp Cys Thr Gly Leu Asn Leu Leu Ser
            20                  25                  30
Ser Ser Asp Arg Pro Thr Ser Ala Ser Pro Val Ala Gly Thr Ile Ala
        35                  40                  45
Val Gln His His Ala Trp Leu Ile Phe  *
    50                  55      57
```

<210> 1999
<211> 66
<212> PRT
<213> Homo sapiens

<400> 1999

```
Met Trp Leu Leu Val Thr Leu Ser Pro Arg Leu Leu Leu Ser Pro Ser
 1               5                   10                  15
His Phe Thr Leu Glu Gly Pro Gln Ile Asp Gln Ala His Ser Glu Leu
            20                  25                  30
Gln Val Leu Pro Leu Val Arg Pro Ser Ala Val Pro Leu Leu Gln Arg
        35                  40                  45
Ala Ser Trp Leu Arg Ser Arg Cys Leu His Leu Pro Lys Thr Val Leu
    50                  55                  60
Val  *
65
```

<210> 2000
<211> 106
<212> PRT
<213> Homo sapiens

<400> 2000

```
Met Gly Arg Cys Leu Ser Leu Gly Ile Leu Arg Gln Gly Leu Cys Cys
 1               5                   10                  15
Pro Cys Trp Ser Val Val Ala Glu Ser Gly Leu Thr Ala Ser Leu Gly
            20                  25                  30
Gly Ser Gly His Pro Ala Thr Ser Cys Ser Lys Glu Ala Gly Thr Thr
        35                  40                  45
Gly Glu Cys Met His His Thr Gln Leu Gly Ile Gln Thr Leu Arg Thr
    50                  55                  60
Tyr Tyr Met Pro Asp Ser Val Glu Leu Ser Glu Thr Met Ser Gly Cys
65                  70                  75                  80
Asn Trp Leu Pro Thr Gln Gln Thr Gln Ser Trp Ala Asn Ile Leu Arg
                85                  90                  95
Val Tyr Leu Thr Leu Lys Tyr Arg Phe Ser
            100                 105 106
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<210> 2001
<211> 88
<212> PRT
<213> Homo sapiens

<400> 2001
Met Glu Arg Arg Arg Leu Leu Gly Gly Met Ala Leu Leu Leu Leu Gln
1 5 10 15
Ala Leu Pro Asn Pro Leu Ser Ala Arg Ala Glu Pro Pro Gln Val Arg
20 25 30
Gly Arg Gly Arg Leu Gly His Val Gly Ser Trp Gly Ser Ser Arg Pro
35 40 45
Gly Trp Arg Gly Leu Lys Glu Cys Cys Cys Gln Glu Leu Arg Gly Pro
50 55 60
Glu Arg Gly Val Tyr Ala Trp Arg Gly Gln Asp Leu Lys Gly Arg Arg
65 70 75 80
Tyr Leu Ala Glu Gly His Leu *
85 87

<210> 2002
<211> 85
<212> PRT
<213> Homo sapiens

<400> 2002
Met Arg Lys Leu Ile Ala Gly Leu Ile Phe Leu Lys Ile Trp Thr Cys
1 5 10 15
Thr Val Arg Thr Ser Thr Asp Leu Pro Gln Thr Glu Asp Cys Ser Gln
20 25 30
Cys Ile His Gln Val Thr Glu Ile Gly Gln Lys Val Ala Thr Val Leu
35 40 45
Leu Phe Tyr Gly Tyr Tyr Lys Tyr Thr Gly Thr Leu Lys Arg Thr Cys
50 55 60
Leu Tyr Asn Val Ile Leu Tyr Lys Val Tyr Ser Pro Gly Asn Asp Gln
65 70 75 80
Pro Asp Val Leu *
84

<210> 2003
<211> 46
<212> PRT
<213> Homo sapiens

<400> 2003
Met Ala Phe Ala Ser Val Leu Leu Ala Arg Ala Ser Pro Ala Val Val
1 5 10 15
Arg Ala Cys Leu Ser Arg Cys Ala Tyr Gly Val Gly Ser Asp Cys Pro
20 25 30
His Leu Val Thr Leu Ala Ala Leu Ile Leu Phe Trp Val *
35 40 45

<210> 2004
<211> 51
<212> PRT
<213> Homo sapiens

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     <400> 2004
Met Trp Leu Phe Ile Ala Ser Lys Cys Ile Phe Leu Leu Ile Val Pro
 1               5                  10                  15
Asn Phe Ile Phe Val Phe Trp Arg Lys Val Phe Ser His Asp Arg Leu
            20                  25                  30
Asn Ile Ala Tyr Ser Phe Glu Leu Ser Ser Lys Tyr Ile Phe Ile Leu
        35                  40                  45
Phe Ile  *
    50


     <210> 2005
     <211> 66
     <212> PRT
     <213> Homo sapiens


     <400> 2005
Met Val Glu Val Val Ser Leu Leu His Leu Tyr Ala Val Ala Cys Ala
 1               5                  10                  15
Arg Lys Gly Pro Phe Pro Asn Thr Lys Asp Leu Ser Gly Trp Thr Pro
            20                  25                  30
Ser Ser Gly Arg Glu Glu Leu Trp Lys Gly Lys Arg Ala Ala Ala Ala
        35                  40                  45
Thr Arg Asn Pro Leu Val Leu Thr Gly Leu Gly Ser Pro Ser Ala Arg
    50                  55                  60
Leu  *
 65


     <210> 2006
     <211> 46
     <212> PRT
     <213> Homo sapiens


     <400> 2006
Met Leu Val Pro Thr Phe Phe Leu Leu Ser Leu Leu Asp Gln Ser Cys
 1               5                  10                  15
Leu Ser Ile Cys Val Ser Gln Asp Tyr Phe Ser Ser Ile Val Val Gln
            20                  25                  30
Ile Arg Gln Ile Gly Ser Leu Cys Leu Asn Lys Ser Leu  *
        35                  40                  45


     <210> 2007
     <211> 87
     <212> PRT
     <213> Homo sapiens


     <400> 2007
Met Pro Thr Leu Ala Lys Trp Ile Leu Ser Leu Ser Met Thr Ser Thr
 1               5                  10                  15
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Thr Trp Ser Pro Cys Ser Ser Met Ile Pro Leu Met Ala Ser Ser Thr
20 25 30
Ala Pro Ser Arg Leu Arg Thr Gly Ser Leu Pro Ser Met Thr Ile Pro
35 40 45
Ser Pro Ser Arg Arg Ser Glu Ile Pro Pro Lys Ser Ser Gly Val Met
50 55 60
Pro Ala Leu Ile Ile Leu Trp Arg Pro Pro Ala Ser Leu Pro Ala Trp
65 70 75 80
Arg Arg Leu Gly Ile Thr *
85 86

<210> 2008
<211> 58
<212> PRT
<213> Homo sapiens

<400> 2008
Met Pro Ala Ile Val Val Phe Leu Phe Cys Phe Val Ile Ser Asp Gly
1 5 10 15
Leu Thr Leu Ser Pro Arg Leu Asp Cys Thr Gly Leu Asn Leu Leu Ser
20 25 30
Ser Ser Asp Arg Pro Thr Ser Ala Ser Pro Val Ala Gly Thr Ile Ala
35 40 45
Val Gln His His Ala Trp Leu Ile Phe *
50 55 57

<210> 2009
<211> 46
<212> PRT
<213> Homo sapiens

<400> 2009
Met Leu Met Tyr Met Phe Tyr Val Leu Pro Phe Cys Gly Leu Ala Ala
1 5 10 15
Tyr Ala Leu Thr Phe Pro Gly Cys Ser Trp Leu Pro Asp Trp Ala Leu
20 25 30
Val Phe Ala Gly Gly Ile Gly Gln Ala Gln Phe Ser His Met
35 40 45 46

<210> 2010
<211> 235
<212> PRT
<213> Homo sapiens

<400> 2010
Met Glu Leu Gly Cys Trp Thr Gln Leu Gly Leu Thr Phe Leu Gln Leu
1 5 10 15
Leu Leu Ile Ser Ser Leu Pro Arg Glu Tyr Thr Val Ile Asn Glu Ala
20 25 30
Cys Pro Gly Ala Glu Trp Asn Ile Met Cys Arg Glu Cys Cys Glu Tyr

```
        35                  40                  45
Asp Gln Ile Glu Cys Val Cys Pro Gly Lys Arg Glu Val Val Gly Tyr
    50                  55                  60
Thr Ile Pro Cys Cys Arg Asn Glu Glu Asn Glu Cys Asp Ser Cys Leu
65                  70                  75                  80
Ile His Pro Gly Cys Thr Ile Phe Glu Asn Cys Lys Ser Cys Arg Asn
                85                  90                  95
Gly Ser Trp Gly Gly Thr Leu Asp Asp Phe Tyr Val Lys Gly Phe Tyr
            100                 105                 110
Cys Ala Glu Cys Arg Ala Gly Trp Tyr Gly Gly Asp Cys Met Arg Cys
        115                 120                 125
Gly Gln Val Leu Arg Ala Pro Lys Gly Gln Ile Leu Leu Glu Ser Tyr
    130                 135                 140
Pro Leu Asn Ala His Cys Glu Trp Thr Ile His Ala Lys Pro Gly Phe
145                 150                 155                 160
Val Ile Gln Leu Arg Phe Val Met Leu Ser Leu Glu Phe Asp Tyr Met
                165                 170                 175
Cys Gln Tyr Asp Tyr Val Glu Gly Cys Asp Gly Asp Asn Arg Asp Gly
            180                 185                 190
His Ile Ile Lys Arg Val Cys Gly Asn Glu Arg Ala Ala Pro Ile His
        195                 200                 205
Asn Ile Arg Ile Leu Thr Ser Arg Pro Phe Pro Leu Pro Gly Leu Ser
    210                 215                 220
Lys Ile Leu Thr Gly Phe His Ala Pro Phe  *
225                 230                 234
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     <210> 2011
     <211> 61
     <212> PRT
     <213> Homo sapiens


     <400> 2011
Met Val Phe Ala Trp Gly Leu Ala Val Asn Lys Thr Ser Leu Val Pro
 1               5                  10                  15
Ile Phe Met Asp Leu Ser Leu Ala Gly Lys Ile Tyr Ile Lys Gln Arg
            20                  25                  30
Met Arg Met Glu Glu Asn Leu Leu Gly Asp Asn Glu Val Lys Glu Glu
        35                  40                  45
Lys Asp Gln Ala Val Lys Trp Gln Thr Leu Arg Trp  *
    50                  55                  60
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     <210> 2012
     <211> 107
     <212> PRT
     <213> Homo sapiens


     <400> 2012
Met Ile Arg Cys Gly Leu Ala Cys Glu Arg Cys Arg Trp Phe Leu Thr
 1               5                  10                  15
Leu Leu Leu Leu Ser Ala Ile Ala Phe Asp Ile Ile Ala Leu Ala Gly
            20                  25                  30
Arg Gly Trp Leu Gln Ser Ser Asp Arg Val Gln Thr Ser Ser Leu Trp
        35                  40                  45
```

Arg Arg Cys Phe Leu Pro Gln Gly Arg Arg Arg Arg Gln Arg Val Leu
50 55 60
Arg Gly Arg Leu Pro Gln Pro His Gly Val Arg Val Gly Ser Ser Ser
65 70 75 80
Ala Ala Met Leu Phe Trp Gly Val Ser Ile Leu Glu Ile Cys Phe Ile
85 90 95
Leu Ser Phe Phe Val Leu Cys Val Pro Gln Ile
100 105 107

<210> 2013
<211> 67
<212> PRT
<213> Homo sapiens

<400> 2013
Met Gly Val Val Leu Tyr Val Leu Val Cys Gly Ala Leu Pro Phe Asp
1 5 10 15
Gly Pro Thr Leu Pro Ile Leu Arg Gln Arg Val Leu Gly Arg Lys Ile
20 25 30
Pro Asp Ser Val Phe His Val Arg Arg Leu Arg Ala Pro Tyr Pro Lys
35 40 45
Asp Val Gly Pro Arg Pro Ile Gln Thr Ala Asn His Ser Pro Asn Gln
50 55 60
Gly Ala *
65 66

<210> 2014
<211> 59
<212> PRT
<213> Homo sapiens

<400> 2014
Met Phe Leu Arg Phe Pro Leu Arg Phe Gly Ile Leu Ala Asp Lys Leu
1 5 10 15
Ile Leu Tyr Lys Ala Ser His Phe Thr Met Leu Ser Val Pro Gly Leu
20 25 30
Tyr Leu Ser Thr Leu Leu Glu Gly Ile Phe Ile Leu Lys Lys Leu Ser
35 40 45
Phe Met Arg Arg Met Gly Val His Ala Thr *
50 55 58

<210> 2015
<211> 55
<212> PRT
<213> Homo sapiens

<400> 2015
Met Val Arg Leu Gln Val Leu Val Leu Val Phe Arg Val Val Gly Ser
1 5 10 15
Gln Gln Met Leu Arg Gln Gly Ala Ala Gly Ala Arg Ser His Arg Val

```
            20                  25                  30
Leu Ala Ser Leu His Phe Gln His Gly Phe Gly Thr Phe His Thr Pro
        35                  40                  45
Ala Arg Ala Gly Gly Ser Glu
    50                  55
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<210> 2016
<211> 64
<212> PRT
<213> Homo sapiens

<400> 2016

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Met Ser Leu Arg Phe Cys Phe Cys Leu Pro Val Cys Pro Ser Leu Pro
 1               5                  10                  15
Ile Ser Val Phe His Val Phe Leu Ser Val Ser Asp His Pro Val Ser
            20                  25                  30
Leu Cys Leu Thr Val Ser Gly His Glu Met Ser Val Ile Val Ala Arg
        35                  40                  45
Phe Thr Leu Ser Leu Tyr Leu Phe Pro Leu Arg Ser Gly Ile Ser  *
    50                  55                  60          63
```

<210> 2017
<211> 58
<212> PRT
<213> Homo sapiens

<400> 2017

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Met Ile Leu Leu Leu Ser Thr Phe Phe Cys Cys Phe Arg Glu Asp Ser
 1               5                  10                  15
Cys Phe Tyr Lys Lys Tyr Val Gly Leu Val Gln Trp Leu Met Pro Val
            20                  25                  30
Ile Pro Ala Leu Trp Glu Ala Lys Val Gly Gly Ser Leu Glu Val Trp
        35                  40                  45
Ser Ser Arg Pro Ala Trp Pro Ile Arg  *
    50                  55      57
```

<210> 2018
<211> 66
<212> PRT
<213> Homo sapiens

<400> 2018

```
Met Leu His Ile Ser Ser Ala Phe His Cys Tyr Ala Phe Leu Pro Leu
 1               5                  10                  15
Phe Ala Leu Thr His Asn Phe Ile Phe Leu Phe Tyr Leu Leu Ser Leu
            20                  25                  30
Ser Pro Lys Leu Glu Cys Lys Phe Gln Glu Gly Arg Asp Phe Tyr Leu
        35                  40                  45
Phe Phe Phe Val Phe Pro Ile Phe Trp His Val Trp His Arg Lys Gly
    50                  55                  60
```

Ile *
65

PATENT COOPERATION TREATY

# PCT

## DECLARATION OF NON-ESTABLISHMENT OF INTERNATIONAL SEARCH REPORT

(PCT Article 17(2)(a), Rule 13*ter*.1(c) and 39)

| Applicant's or agent's file reference<br>21272-018 | **IMPORTANT DECLARATION** | Date of mailing (day/month/year)<br>27 JUN 2001 |
|---|---|---|
| International application No.<br>PCT/US01/02687 | International filing date *(day/month/year)*<br>25 January 2001 (25.01.2001) | (Earliest) Priority date *(day/month/year)*<br>25 January 2000 (25.01.2000) |
| International Patent Classification (IPC) or both national classification and IPC<br>IPC(7): C12P 21/06 and US Cl.: 435/69.1 | | |
| Applicant<br>HYSEQ, INC. | | |

This International Searching Authority hereby declares, according to Article 17(2)(a), that **no international search report will be established** on the international application for the reasons indicated below.

1. ☐ The subject matter of the international application relates to:
   a. ☐ scientific theories.
   b. ☐ mathematical theories
   c. ☐ plant varieties.
   d. ☐ animal varieties.
   e. ☐ essential biological processes for the production of plants and animals, other than microbiological processes and the products of such processes.
   f. ☐ schemes, rules or methods of doing business.
   g. ☐ schemes, rules or methods of performing purely mental acts.
   h. ☐ schemes, rules or methods of playing games.
   i. ☐ methods for treatment of the human body by surgery or therapy.
   j. ☐ methods for treatment of the animal body by surgery or therapy.
   k. ☐ diagnostic methods practised on the human or animal body.
   l. ☐ mere presentations of information.
   m. ☐ computer programs for which this International Searching Authority is not equipped to search prior art.

2. ☒ The failure of the following parts of the international application to comply with prescribed requirements prevents a meaningful search from being carried out:
   ☐ the description ☒ the claims ☐ the drawings

3. ☒ The failure of the nucleotide and/or amino acid sequence listing to comply with the standard provided for in Annex C of the Administrative Instructions prevents a meaningful search from being carried out:
   ☐ the written form has not been furnished or does not comply with the standard.
   ☒ the computer readable form has not been furnished or does not comply with the standard.

4. Further comments:

| Name and mailing address of the ISA/US<br>Commissioner of Patents and Trademarks<br>Box PCT<br>Washington, D.C. 20231 | Authorized officer<br>Young J. Kim |
|---|---|